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Lacourciere, Karen

Sent:

Thursday, January 09, 2003 12:38 PM STIC-Biotech/ChemLib

To:

Subject:

sequence search request 09/757,100

Please search SEQ ID NO:18 of 09/757,100 in both the commercial databases and the pending files (Interference)

Thank-you!--

Karen A. Lacourciere Ph.D.

CM1 11D09 GAU 1635

(703) 208-7523 mailbox 11E12

Point of Contact: Beverly Shears Technical Info. Specialist CM1 1E05 Tel: 308-4994

Searcher: Phone: Location: Date Picked Up:____ Date Completed: Searcher Prep/Review:____ Clerical:_____ Online time:

TYPE OF SEARCH: NA Sequences:_____ AA Sequences:_____ Structures: Bibliographic:_____ Litigation:_____ Full text:____ Patent Family: Other:____

VENDOR/COST (where applic.) STN:_____ DIALOG:____ Questel/Orbit: DRLink:__ Lexis/Nexis:___ Sequence Sys.:_____ WWW/Internet:_____ Other (specify):__

U.S. DEPARTMENT OF COMMERCE Patent and Trademark Office

SEARCH REQUEST FORM

Requestor's Name:	Serial Number:	
Date:	Phone:	Art Unit:
Search Topic: Please write a detailed statement of search topic. that may have a special meaning. Give examples a copy of the sequence. You may include a cop	or relevant citations, authors keywords.	etc., if known. For sequences, please attach
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Date completed: 01-13-03 Searcher: Belevily e 4:17 Terminal time: 23 Elapsed time: CPU time: Total time:	Search Site STIC CM-1 Pre-S Type of Search N.A. Seque	
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(FILE 'REGISTRY' ENTERED AT 10:55:53 ON 13 JAN 2003)
L1 20 S TTTCAACCAGATGGTCATTC/SQSN

FILE 'HCAPLUS' ENTERED AT 10:57:04 ON 13 JAN 2003 7 S L1

L2 ANSWER 1 OF 7 HCAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2002:889441 HCAPLUS

DOCUMENT NUMBER: 137:381006

TITLE: Expressed sequence profiles and their use for

the therapy and diagnosis of human lung cancer Henderson, Robert A.; Wang, Tongtong; Watanabe, Yoshihiro; Johnson, Jeffrey C.; Retter, Marc W.; Durham, Margarita; Carter, Darrick; Fanger, Gary

R.; Vedvick, Thomas S.; Bangur, Chaitanya S.;

McNabb, Andria

PATENT ASSIGNEE(S): Corixa Corporation, USA

SOURCE: U.S. Pat. Appl. Publ., 82 pp., Cont.-in-part of

U.S. Ser. No. 849,626.

CODEN: USXXCO

DOCUMENT TYPE: Patent

LANGUAGE: English FAMILY ACC. NUM. COUNT: 5

PATENT INFORMATION:

INVENTOR(S):

	PATENT NO.	KIND	DATE		APPLICATION NO).	DATE
PRIO	PATENT NO. US 2002172952 US 6504010 US 2002168637 US 2002197669 RITY APPLN. INFO.	A1 B1 A1 A1	DATE 20021121 20030107 20021114 20021226	US US US US US US US US US US US US	US 2001-902943 US 2000-702703 US 2000-736453 US 2001-849626 1999-419356 1999-466867 1999-476300 2000-519642 2000-533077 2000-546259 2000-560406 2000-589184 2000-651563 2000-651563 2000-658824 2000-677419	A2 A2 A2 A2 A2 A2 A2 A2 A2 A2 A2 A2	20010710 20001030 20001213 20010503 19990630 19991015 19991217 19991230 20000306 20000322 20000410 20000427 20000605 20000711 20000829 20000908 20000926 20001006
				US US US	2000-677419 2000-702705 2000-736457 2001-849626	A2 A2 A2 A2	20001006 20001030 20001213 20010503
				WO	2000-US18061	Α2	20000630

AB Nucleic acid and encoded protein compns. and methods for the therapy and diagnosis of cancer, particularly lung cancer, are disclosed. Thus, cDNAs encoding tumor-assocd. proteins were identified by conventional and PCR-based library subtraction, cDNA microarray anal., and real-time PCR using cDNA libraries isolated from lung adenocarcinoma, small cell lung carcinoma, squamous cell lung carcinoma, and lung neuroendocrine carcinoma vs. a pool of poly(A)+RNA from 9 normal tissues. Illustrative compns. comprise one or more lung tumor polypeptides, immunogenic portions thereof,

polynucleotides that encode such polypeptides, antigen presenting cell that expresses such polypeptides, and T cells that are specific for cells expressing such polypeptides. The disclosed compns. are useful, for example, in the diagnosis, prevention and/or treatment of diseases, particularly lung cancer.

IT 475864-34-7

RL: ANT (Analyte); BSU (Biological study, unclassified); DGN (Diagnostic use); PRP (Properties); THU (Therapeutic use); ANST (Analytical study); BIOL (Biological study); USES (Uses) (nucleotide sequence; expressed sequence profiles and their use for the therapy and diagnosis of human lung cancer)

L2 ANSWER 2 OF 7 HCAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2002:869469 HCAPLUS

DOCUMENT NUMBER: 137:364446

TITLE: Expressed sequence profiles and their use for the therapy and diagnosis of human lung cancer

INVENTOR(S): Wang, Tongtong; Bangur, Chaitanya S.; Lodes, Michael J.; Fanger, Gary R.; Vedvick, Thomas S.;

Carter, Darrick; Retter, Marc W.; Mannion, Jane;

Fan, Liqun; Wang, Aijun

PATENT ASSIGNEE(S): USA

SOURCE: U.S. Pat. Appl. Publ., 66 pp., Cont.-in-part of

U. S. Ser. No. 702,705.

CODEN: USXXCO

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 5

PATENT INFORMATION:

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                                                          DATE
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US 2002168637
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        TJ, TM
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                CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
EP 1196565
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                                        EP 2000-945031
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US 2002197669
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US 2002172952
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WO 2002004514
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        NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR,
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TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ,
             MD, RU, TJ, TM
         RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH,
             CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE,
             TR, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD,
             TG
PRIORITY APPLN. INFO .:
                                        US 1999-346492
                                                          A2 19990630
                                        US 1999-419356
                                                          A2 19991015
                                        US 1999-466867
                                                          A2 19991217
                                        US 1999-476300
                                                          A2 19991230
                                        US 2000-519642
                                                          A2 20000306
                                        US 2000-533077
                                                          A2 20000322
                                        US 2000-546259
                                                          A2 20000410
                                                          A2 20000427
                                        US 2000-560406
                                        US 2000-589184
                                                          A2 20000605
                                        WO 2000-US18061
                                                         A2 20000630
                                        US 2000-614124
                                                          A2 20000711
                                        US 2000-651563
                                                          A2 20000829
                                        US 2000-658824
                                                          A2 20000908
                                                         A2 20000926
                                        US 2000-671325
                                        US 2000-677419
                                                          A2 20001006
                                        US 2000-702705
                                                          A2 20001030
                                        US 2000-736457
                                                          A2 20001213
                                        US 2001-849626
                                                          A2 20010503
     Nucleic acid and encoded protein compns. and methods for the therapy
AΒ
     and diagnosis of cancer, particularly lung cancer, are disclosed.
     Thus, cDNAs encoding tumor-assocd. proteins were identified by
     conventional and PCR-based library subtraction, cDNA microarray
     anal., and real-time PCR using cDNA libraries isolated from lung
     adenocarcinoma, small cell lung carcinoma, squamous cell lung
     carcinoma, and lung neuroendocrine carcinoma vs. a pool of poly(A)+
     RNA from 9 normal tissues. Illustrative compns. comprise one or
     more lung tumor polypeptides, immunogenic portions thereof,
     polynucleotides that encode such polypeptides, antigen presenting
     cell that expresses such polypeptides, and T cells that are specific
     for cells expressing such polypeptides. The disclosed compns. are
     useful, for example, in the diagnosis, prevention and/or treatment
     of diseases, particularly lung cancer.
IT
     475019-40-0
     RL: ANT (Analyte); BSU (Biological study, unclassified); DGN
     (Diagnostic use); PRP (Properties); THU (Therapeutic use); ANST
     (Analytical study); BIOL (Biological study); USES (Uses)
        (nucleotide sequence; expressed sequence profiles and their use
        for the therapy and diagnosis of human lung cancer)
L2
     ANSWER 3 OF 7 HCAPLUS COPYRIGHT 2003 ACS
                         2002:332807 HCAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                         136:351185
TITLE:
                         The contribution of 700,000 ORF sequence tags to
                         the definition of the human transcriptome
AUTHOR(S):
                         Camargo, Anamaria A.; Samaia, Helena P. B.;
                         Dias-Neto, Emmanuel; Simao, Daniel F.; Migotto,
                         Italo A.; Briones, Marcelo R. S.; Costa,
                         Fernando F.; Nagai, Maria Aparecida;
                         Verjovski-Almeida, Sergio; Zago, Marco A.;
                         Andrade, Luis Eduardo C.; Carrer, Helaine;
                         El-Dorry, Hamza F. A.; Espreafico, Enilza M.;
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Searcher: Shears 308-4994

Habr-Gama, Angelita; Giannella-Neto, Daniel;

Goldman, Gustavo H.; Gruber, Arthur; Hackel, Christine; Kimura, Edna T.; Maciel, Rui M. B.; Marie, Suely K. N.; Martins, Elizabeth A. L.; Nobrega, Marina P.; Paco-Larson, Maria Luisa; Pardini, Maria Ines M. C.; Pereira, Goncalo G.; Pesquero, Joao Bosco; Rodrigues, Vanderlei; Rogatto, Silvia R.; Da Silva, Ismael D. C. G.; Sogayar, Mari C.; Sonati, Maria De Fatima; Tajara, Eloiza H.; Valentini, Sandro R.; Alberto, Fernando L.; Amaral, Maria Elisabete J.; Aneas, Ivy; Arnaldi, Liliane A. T.; De Assis, Angela M.; Bengtson, Mario Henrique; Bergamo, Nadia Aparecida; Bombonato, Vanessa; De Camargo, Maria E. R.; Canevari, Renata A.; Carraro, Dirce M.; Cerutti, Janete M.; Correa, Maria Lucia C.; Correa, Rosana F. R.; Costa, Maria Cristina R.; Curcio, Cyntia; Hokama, Paula O. M.; Ferreira, Ari J. S.; Furuzawa, Gilberto K.; Gushiken, Tsieko; Ho, Paulo L.; Kimura, Elza; Krieger, Jose E.; Leite, Luciana C. C.; Majumder, Paromita; Marins, Mozart; Marques, Everaldo R.; Melo, Analy S. A.; Melo, Monica; Mestriner, Carlos Alberto; Miracca, Elisabete C.; Miranda, Daniela C.; Nascimento, Ana Lucia T. O.; Nobrega, Francisco G.; Ojopi, Elida P. B.; Pandolfi, Jose Rodrigo C.; Pessoa, Luciana G.; Prevedel, Aline C.; Rahal, Paula; Rainho, Claudia A.; Reis, Eduardo M. R.; Ribeiro, Marcelo L.; Da Ros, Nancy; De Sa, Renata G.; Sales, Magaly M.; Sant'anna, Simone Cristina; Dos Santos, Mariana L.; Da Silva, Aline M.; Da Silva, Neusa P.; Silva, Wilson A., Jr.; Da Silveira, Rosana A.; Sousa, Josane F.; Stecconi, Daniella; Tsukumo, Fernando; Valente, Valeria; Soares, Fernando; Moreira, Eloisa S.; Nunes, Diana N.; Correa, Ricardo G.; Zalcberg, Heloisa; Carvalho, Alex F.; Reis, Luis F. L.; Brentani, Ricardo R.; Simpson, Andrew J. G.; De Souza, Sandro J.

CORPORATE SOURCE:

Ludwig Institute for Cancer Research, Sao Paulo,

01509-010, Brazil

Proceedings of the National Academy of Sciences SOURCE: of the United States of America (2001), 98(21),

12103-12108

CODEN: PNASA6; ISSN: 0027-8424 National Academy of Sciences

PUBLISHER: DOCUMENT TYPE:

Journal LANGUAGE: English

AΒ Open reading frame expressed sequences tags (ORESTES) differ from conventional ESTs by providing sequence data from the central protein coding portion of transcripts. A total of 696,745 ORESTES sequences were generated from 24 human tissues and a subset of the data that correspond to a set of 15,095 full-length mRNAs used as a means of assessing the efficiency of the strategy and its potential contribution to the definition of the human transcriptome. It was estd. that ORESTES sampled over 80% of all highly and moderately expressed, and between 40% and 50% of rarely expressed, human genes. In the most thoroughly sequenced tissue, the breast, the 130,000

ORESTES generated are derived from transcripts from an estd. 70% of all genes expressed in that tissue, with an equally efficient representation of both highly and poorly expressed genes. respect, the capacity of the ORESTES strategy both for gene discovery and shotgun transcript sequence generation significantly exceeds that of conventional ESTs. The distribution of ORESTES is such that many human transcripts are now represented by a scaffold of partial sequences distributed along the length of each gene product. The exptl. joining of the scaffold components, by reverse transcription-PCR, represents a direct route to transcript finishing that may represent a useful alternative to full-length cDNA cloning. [This abstr. record is one of many records for this document necessitated by the large no. of index entries required to fully index the document and publication system constraints.].

IT 270220-29-6, GenBank AW992843 270220-33-2, GenBank AW992847

RL: BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study)

(nucleotide sequence; contribution of 700,000 ORF sequence tags to the definition of the human transcriptome)

ANSWER 4 OF 7 HCAPLUS COPYRIGHT 2003 ACS L2 ACCESSION NUMBER: 2002:332795 HCAPLUS 136:351174

DOCUMENT NUMBER:

TITLE:

AUTHOR(S):

the definition of the human transcriptome Camargo, Anamaria A.; Samaia, Helena P. B.; Dias-Neto, Emmanuel; Simao, Daniel F.; Migotto, Italo A.; Briones, Marcelo R. S.; Costa, Fernando F.; Nagai, Maria Aparecida; Verjovski-Almeida, Sergio; Zago, Marco A.; Andrade, Luis Eduardo C.; Carrer, Helaine; El-Dorry, Hamza F. A.; Espreafico, Enilza M.; Habr-Gama, Angelita; Giannella-Neto, Daniel; Goldman, Gustavo H.; Gruber, Arthur; Hackel, Christine; Kimura, Edna T.; Maciel, Rui M. B.; Marie, Suely K. N.; Martins, Elizabeth A. L.; Nobrega, Marina P.; Paco-Larson, Maria Luisa; Pardini, Maria Ines M. C.; Pereira, Goncalo G.; Pesquero, Joao Bosco; Rodrigues, Vanderlei; Rogatto, Silvia R.; Da Silva, Ismael D. C. G.; Sogayar, Mari C.; Sonati, Maria De Fatima; Tajara, Eloiza H.; Valentini, Sandro R.; Alberto, Fernando L.; Amaral, Maria Elisabete

The contribution of 700,000 ORF sequence tags to

J.; Aneas, Ivy; Arnaldi, Liliane A. T.; De Assis, Angela M.; Bengtson, Mario Henrique; Bergamo, Nadia Aparecida; Bombonato, Vanessa; De Camargo, Maria E. R.; Canevari, Renata A.; Carraro, Dirce M.; Cerutti, Janete M.; Correa, Maria Lucia C.; Correa, Rosana F. R.; Costa, Maria Cristina R.; Curcio, Cyntia; Hokama, Paula O. M.; Ferreira, Ari J. S.; Furuzawa, Gilberto K.; Gushiken, Tsieko; Ho, Paulo L.; Kimura, Elza; Krieger, Jose E.; Leite, Luciana C. C.; Majumder, Paromita; Marins, Mozart; Marques, Everaldo R.; Melo, Analy S. A.; Melo, Monica;

Mestriner, Carlos Alberto; Miracca, Elisabete C.; Miranda, Daniela C.; Nascimento, Ana Lucia



T. O.; Nobrega, Francisco G.; Ojopi, Elida P. B.; Pandolfi, Jose Rodrigo C.; Pessoa, Luciana G.; Prevedel, Aline C.; Rahal, Paula; Rainho, Claudia A.; Reis, Eduardo M. R.; Ribeiro, Marcelo L.; Da Ros, Nancy; De Sa, Renata G.; Sales, Magaly M.; Sant'anna, Simone Cristina; Dos Santos, Mariana L.; Da Silva, Aline M.; Da Silva, Neusa P.; Silva, Wilson A., Jr.; Da Silveira, Rosana A.; Sousa, Josane F.; Stecconi, Daniella; Tsukumo, Fernando; Valente, Valeria; Soares, Fernando; Moreira, Eloisa S.; Nunes, Diana N.; Correa, Ricardo G.; Zalcberg, Heloisa; Carvalho, Alex F.; Reis, Luis F. L.; Brentani, Ricardo R.; Simpson, Andrew J. G.; De Souza, Sandro J.

CORPORATE SOURCE:

Ludwig Institute for Cancer Research, Sao Paulo,

01509-010, Brazil

SOURCE:

Proceedings of the National Academy of Sciences of the United States of America (2001), 98(21),

12103-12108

CODEN: PNASA6; ISSN: 0027-8424 National Academy of Sciences

PUBLISHER:
DOCUMENT TYPE:

Journal English

LANGUAGE: Open reading frame expressed sequences tags (ORESTES) differ from AB conventional ESTs by providing sequence data from the central protein coding portion of transcripts. A total of 696,745 ORESTES sequences were generated from 24 human tissues and a subset of the data that correspond to a set of 15,095 full-length mRNAs used as a means of assessing the efficiency of the strategy and its potential contribution to the definition of the human transcriptome. It was estd. that ORESTES sampled over 80% of all highly and moderately expressed, and between 40% and 50% of rarely expressed, human genes. In the most thoroughly sequenced tissue, the breast, the 130,000 ORESTES generated are derived from transcripts from an estd. 70% of all genes expressed in that tissue, with an equally efficient representation of both highly and poorly expressed genes. In this respect, the capacity of the ORESTES strategy both for gene discovery and shotgun transcript sequence generation significantly exceeds that of conventional ESTs. The distribution of ORESTES is such that many human transcripts are now represented by a scaffold of partial sequences distributed along the length of each gene product. The exptl. joining of the scaffold components, by reverse transcription-PCR, represents a direct route to transcript finishing that may represent a useful alternative to full-length cDNA cloning. [This abstr. record is one of many records for this document necessitated by the large no. of index entries required to fully index the document and publication system constraints.].

IT **272727-43-2**, GenBank BE092864

RL: BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study)

(nucleotide sequence; contribution of 700,000 ORF sequence tags to the definition of the human transcriptome)

L2 ANSWER 5 OF 7 HCAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2002:51525 HCAPLUS

DOCUMENT NUMBER:

136:129944

TITLE:

Human nucleic acid and expressed protein

IIIIO FAUL DEMINA (USPTU)

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compositions and methods for the therapy and
                           diagnosis of lung cancer
INVENTOR(S):
                           Wang, Tongtong; Watanabe, Yoshihiro; Henderson,
                           Robert A.; Johnson, Jeffrey C.; Retter, Marc W.;
                           Marnerakis, Margarita; Carter, Darrick; Fanger,
                           Gary Richard; Vedvick, Thomas S.; Bangur,
                           Chaitanya S.; McNabb, Andria; Wang, Aijun;
                           Fanger, Neil; Switzer, Ann; McNeill, Patricia
                           D.; Clapper, Jonathan D.
                           Corixa Corporation, USA
PATENT ASSIGNEE(S):
SOURCE:
                           PCT Int. Appl., 223 pp.
                           CODEN: PIXXD2
DOCUMENT TYPE:
                           Patent
LANGUAGE:
                           English
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
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                                              APPLICATION NO.
                                                                DATE
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                                           US 2000-589184
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                                           WO 2000-US18061 A2 20000630
AΒ
     Compns. and methods for the therapy and diagnosis of cancer,
     particularly lung cancer, are disclosed. Illustrative compns.
     comprise one or more lung tumor polypeptides, immunogenic portions
     thereof, the cDNA polynucleotides that encode such polypeptides,
     antigen-presenting cell that expresses such polypeptides, and T
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Searcher: Shears 308-4994

cells that are specific for cells expressing such polypeptides.

Thus, lung tumor cDNAs are isolated and characterized using conventional cDNA library subtraction, PCR-based subtraction, microarray anal., and lung chip 5. The cDNA libraries are isolated from lung adenocarcinoma, small cell lung carcinoma, lung neuroendocrine carcinoma, and squamous cell lung carcinoma. The immunogenic gene products are expressed in Escherichia coli, and epitope mapping is provided using antibodies one of the lung tumor-specific antigens, designated L552S. The disclosed compns. are useful, for example, in the diagnosis, prevention and/or treatment of diseases, particularly lung cancer.

IT 388152-61-2P

RL: ANT (Analyte); BPN (Biosynthetic preparation); DGN (Diagnostic use); PRP (Properties); THU (Therapeutic use); ANST (Analytical study); BIOL (Biological study); PREP (Preparation); USES (Uses) (nucleotide sequence; human nucleic acid and expressed protein compns. and methods for the therapy and diagnosis of lung cancer)

L2 ANSWER 6 OF 7 HCAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2001:731174 HCAPLUS

DOCUMENT NUMBER: 135:283933

TITLE: Nucleic acid markers and methods for the

measurement of sensitivity or resistance to agents used in identification, assessment, prevention, and therapy of human cancers

INVENTOR(S): Lillie, James; Brown, Jeffrey L.; Bolt, Andrew;

Van Huffel, Christophe

PATENT ASSIGNEE(S): Millennium Predictive Medicine, Inc., USA

SOURCE: PCT Int. Appl., 223 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

PATENT INFORMATION:

PATENT NO.			KI	CIND DATE			APPLICATION NO.					DATE				
	2001								M	0 20	01-U	S920	2	2001	0322	
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US PRIORIT	2002 Y APP	0815			1	2002	0627		US 2	S 20 000- 000-	1921	00P	P	2001 2000 2000	0324	

AB The present invention is directed to the identification of markers that can be used to det. whether cancer cells are sensitive or resistant to a therapeutic agent. The present invention is also directed to the identification of therapeutic targets. The invention features 436 "sensitivity markers". These are markers that are expressed in most or all cell lines that are sensitive to

treatment with an agent and which are not expressed (or are expressed at a rather low level) in cells that are resistant to treatment with that agent. The invention also features 3132 "resistance markers". These are markers that are expressed in most or all cell lines that are resistant to treatment with an agent and which are not expressed (or are expressed at a rather low level) in cells that are sensitive to treatment with that agent. PCR-based subtractive hybridization of cDNA libraries from normal and cancerous tissues was used to identify the markers. At least some of the examples relate to sensitivity or resistance to taxol.

IT 191206-39-0

RL: ANT (Analyte); BOC (Biological occurrence); BSU (Biological study, unclassified); PRP (Properties); THU (Therapeutic use); ANST (Analytical study); BIOL (Biological study); OCCU (Occurrence); USES (Uses)

(resistance marker; nucleic acid markers and methods for the measurement of sensitivity or resistance to agents used in identification, assessment, prevention, and therapy of human cancers)

L2 ANSWER 7 OF 7 HCAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2001:551387 HCAPLUS

DOCUMENT NUMBER: 135:103496

TITLE: Nucleic acid compositions, kits, and methods for

identification, assessment, prevention, and

therapy of human breast cancer

INVENTOR(S): Lillie, James; Palermo, Adam; Wang, Youzhen;

Steinmann, Kathleen; Elias, Josh

PATENT ASSIGNEE(S): Millennium Predictive Medicine, Inc., USA

SOURCE: PCT Int. Appl., 2674 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

PATENT INFORMATION:

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KIND DATE ,
     PATENT NO.
                                                     APPLICATION NO.
     WO 2001046697 A2
                                   20010628
                                                   WO 2000-US35214 20001221
         AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN,
          CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU,
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                                                      US 1999-PV171406 19991221
PRIORITY APPLN. INFO.:
                                                      US 2000-PV176423 20000114
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                                                      US 2000-PV205231 20000515
                                                      US 2000-PV213236 20000620
                                                      US 2000-PV219865 20000720
     The invention relates to nucleic acid marker compns., kits and
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AB The invention relates to nucleic acid marker compns., kits and methods for detecting, characterizing, preventing, and treating human breast cancers. A variety of markers are provided, wherein changes in the levels of expression of one or more of the nucleic acid markers is correlated with the presence of breast cancer. The

level of expression of numerous potential markers was measured in cells obtained from breast cancer tissue samples obtained form fifteen patients afflicted with breast cancer and from eleven breast cancer cell cultures, based on comparison with expression levels of each marker in corresponding non-cancerous breast tissue and cell cultures. The 15 cancer tissue samples include (i) five invasive lobular carcinomas (ILC), (ii) five invasive ductal carcinomas (IDC), and (iii) five samples of ductal carcinoma in situ (DCIS). As an addnl. evaluation of ability to indicate breast cancer, individual markers that were identified by transcriptional profiling criteria were also tested in six different subtracted library expts. In addn., protein profiling expts. were undertaken to assess whether the proteins assocd. with the expression of individual markers of the invention are secreted. Table 21 lists approx. 43,500 GenBank Accession Nos. from the present invention. [This abstr. record is two of eight records for this document necessitated by the large no. of index entries required to fully index the document and publication system constraints.].

IT 191206-39-0

RL: BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study)

(nucleotide sequence; nucleic acid compns., kits, and methods for identification, assessment, prevention, and therapy of human breast cancer)

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L3 ANSWER 1 OF 7 REGISTRY COPYRIGHT 2003 ACS

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CN 981: PN: US20020172952 SEQID: 981 claimed DNA

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L3 ANSWER 2 OF 7 REGISTRY COPYRIGHT 2003 ACS

RN 475019-40-0 REGISTRY

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REFERENCE 1: 137:364446

L3 ANSWER 3 OF 7 REGISTRY COPYRIGHT 2003 ACS

RN **388152-61-2** REGISTRY



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     ANSWER 6 OF 7 REGISTRY COPYRIGHT 2003 ACS
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     4606: PN: WO0146697 TABLE: 21 claimed DNA
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REFERENCE 2: 135:103496

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SUMMARIES

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CURRENT FILING DATE: 2000-07-13
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PRIOR FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: PatentIN Ver: 2.0
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LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase TITLE OF INVENTION: Expression FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: US/09/377,310A
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Gaarde, William A.
Nero, Pamela S.
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1513, Ap
159, App
159, App
16983, Ap
6983, Ap
6983, Ap
6983, App
16983, App
1798, App
1898, App
18
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Sequence 18, Application US/09757100B

| GENERAL INFORMATION:
| APPLICANT: Monia, Brett P. |
| APPLICANT: Morio, William A. |
| APPLICANT: Nero, Pamela S. |
| TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase |
| TITLE OF INVENTION: Expression |
| FILE REFERENCE: ISPH-0533 |
| CURRENT APPLICATION NUMBER: US/09/757,100B |
| CURRENT APPLICATION NUMBER: 09/377,310 |
| PRIOR APPLICATION NUMBER: 09/377,310 |
| PRIOR APPLICATION NUMBER: PCT/US00/18999 |
| PRIOR APPLICATION NUMBER: PCT/US00/18999 |
| PRIOR APPLICATION NUMBER: WOT/US00/18999 |
| PRIOR FILING DATE: 2000-07-13 |
| NUMBER OF SEQ ID NOS: 44 |
| SOFTWARE: Patentin Ver. 2.0 |
| SEQ ID NO 18 |
| ELECTH: 20
                                  Sequence 6610, Application US/09332782
(GENERAL INFORMATION:
APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-754C01
CURRENT APPLICATION NUMBER: US/09/332,782
CURRENT FILING DATE: 1999-06-14
EARLIER APPLICATION NUMBER: US 09/181,430
EARLIER APPLICATION NUMBER: US 09/181,430
EARLIER FILING DATE: 1998-10-28
NUMBER OF SEQ ID NOS: 21027
SOFTWARE: FastSEQ for Windows Version 3.0
LENGTH; 325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 18
LENGTH: 20
TYPE: DNA
TYPE: DNA
FEARTHER.
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; OTHER INFORMATION: antisense sequence
US-09-377-310-18
                                                                                                                                                                                                                                                                                                                                 US-09-332-782-6610/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: antisense sequence US-09-757-100B-18
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Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
ORGANISM: Homo sapiens
                       TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
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20; Conserv
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llarity 100.0%;
Conservative 0
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Pred. No. 5
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TITLE OF INVENTION: GENE EXPRESSION PROFILING OF PRIMARY BREAST CARCINOMAS TITLE OF INVENTION: USING ARRAYS OF CANDIDATE GENES
FILE REPERBRUE: 1546.R-00
CURRENT APPLICATION NUMBER: US/10/007,926A
CURRENT FILING DATE: 2001-12-07
PRIOR APPLICATION NUMBER: 60/254,090
PRIOR PILING DATE: 2000-12-08
NUMBER OF SEQ ID NOS: 468
SOFTWARE: PATENTIN VET. 2.1
        Gaps
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ORGANISM: Homo sapiens
PEATURE:
OTHER INFORMATION: 5' terminal sequence. ptk2 protein tyrosine
OTHER INFORMATION: kinase 2 (PTK2) gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 20; DB 17; Length 379; 100.0%; Pred. No. 9.8; cive 0; Mismatches 0; Indels
                                                                                                                                     RESULT 6

US-09-359-067-23180/C

Sequence 23180, Application US/09359067

Sequence 23180, Application US/09359067

GENERAL INFORMATION:

APPLICANT: Hyseq, Inc.

TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED

TITLE OF INVENTION: PROM VARIOUS CDNA LIBRARIES

FILE REFERENCE: 20411-747CON1

CURRENT APPLICATION NUMBER: US 09/131,598

EARLIER PILING DATE: 1999-07-22

BARLIER FILING DATE: 1998-08-10

NUMBER OF SEQ ID NOS: 49786

SEQ ID NO 2180

SEQ ID NO 2180
      Indels
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      Mismatches
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APPLICANT: BERTUCCI, FRANCOIS
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BIRNBAUM, DANIEL
NGUYEN, CATHERINE
VIENS, PATRICE
FERT, VINCENT
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LOCATION: (275)
OTHER INFORMATION: a, t, c
                                             TTTCAACCAGATGGTCATTC
                                                                                   TTTCAACCAGATGGTCATTC
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Best Local Similarity 100.
Matches 20; Conservative
      Conservative
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OTHER INFORMATION: a, t,
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LOCATION: (364)
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CORGANISM: Homo sapiens
US-09-359-067-23180
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US-10-007-926A-284/c
    20;
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APPLICANT:
APPLICANT:
APPLICANT:
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                                         Score 20; DB 17; Length 325;
Pred. No. 9.5;
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Pred. No. 9.5;
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JITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 754CIP
CURRENT FILING DATE: 2000-02-29
EARLIER PELICATION NUMBER: 09/332,782
EARLIER PELING DATE: 1999-06-14
EARLIER PILING DATE: 1998-10-28
NUMBER OF SEQ ID NOS: 21027
                                                                                   Indels
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                                                                                 Mismatches
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                                       Query Match 100.0%; S
Best Local Similarity 100.0%; P
Matches 20; Conservative 0;
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100.0%;
                                                                                                                                                            53 Trrcaaccacarcarrc 34
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APPLICANT: Asghari, Vida
APPLICANT: Blun, Linda
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Sahourieh, Hannah
Sidhu, Navjiwan
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Engleman, Carrie
Faulkner, Brandy
Fox, Melvin
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Lomelli, Michelle
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APPLICANT: Damavandi, Simin
APPLICANT: Dickson, Mark
APPLICANT: Drake, Jim
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Smythe, Ashleigh
Tkach, Joe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Garcia, Veronica
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kofler, Janette
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Phuong
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Verna, Ron
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ORGANISM: Homo sapiens
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Best Local Similarity
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US-09-332-782-6610
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US-09-824-557-1277/c
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                                                                                     US-09-790-435-4313
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1277
ELOUTH: 421
TYPE: DNA
ORGANISM: Homo sapiens
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Query Match
Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                                 SEQ ID NO 4313
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Best Local Similarity 100.0%;
Matches 20; Conservative 0;
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NAME/KEY: misc feature
LOCATION: (1) ... (421)
OTHER INFORMATION: n = A,T,C or
-09-824-557-1277
                                                                                                                                                                                                                                                                                                                   APPLICANT: Gearing, David P.
TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
TITLE OF INVENTION: THEREFOR
FILE REFERENCE: 1600.2060-001
CURRENT APPLICATION NUMBER: US/09/790,435
CURRENT FILING DATE: 2001-02-22
CURRENT FILING DATE: 2001-02-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Shyjan, Andrew W.
TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
TITLE OF INVENTION: THEREFOR
FILE REFERNCE: 1600.2086-001
CURRENT APPLICATION NUMBER: US/09/824,557
CURRENT FILING DATE: 2001-04-02
PRIOR APPLICATION NUMBER: US 60/193,438
PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 9773
                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 7473
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/183,730 PRIOR FILING DATE: 2000-02-22
                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Holtzman, Douglas A. APPLICANT: Gearing, David P.
                                                                                             FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(433)
OTHER INFORMATION: n = A,T,C or
                                                                                                                                                                      LENGTH: 433
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         270 TTTCAACCAGATGGTCATTC 251
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                             100.0%;
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                           Score 20; DB
Pred. No. 10;
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Pred. No. 10;
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              Mismatches
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                                                Length 433;
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; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (1)... (436)
; OTHER INFORMATION: n = A,T,C or G
US-09-289-768-4292
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US-09-289-768-4292/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Hyseq, Inc.
APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: ROWEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: ROW VARIOUS cDNA LIBRARIES
FILE REFERENCE: 20411-765
CURRENT EDPLICATION NUMBER: US/09/289,768
CURRENT FILING DATE: 1999-04-08
NUMBER OF SEQ ID NOS: 39996
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 4292
                                                                                                                                                                                                                                                                                                                                              Sequence 4292, Application US/09939397
GENERAL INFORMATION:
APPLICANT: HYSEQ, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS CONA LIBRARIES
FILE REFERENCE: 20411-765
CURRENT APPLICATION NUMBER: US/09/939,397
CURRENT APPLICATION NUMBER: US/09/939,397
CURRENT APPLICATION NUMBER: 09/289,768
PRIOR APPLICATION NUMBER: 09/289,768
PRIOR FILING DATE: 1999-04-08
PRIOR FILING DATE: 1999-04-08
                                                                       Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 39996
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 4292
                                                                                                                                                                TYPE: DNA
ORCANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)...(436)
OTHER INFORMATION: n = A,T,C o:
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TYPE: DNA
ORGANISM: Homo sapiens
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Local Similarity 100.0%;
nes 20; Conservative 0
97 TTTCAACCAGATGGTCATTC 78
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                         TTTCAACCAGATGGTCATTC 20
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Pred. No. 1
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Pred. No.
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                                                                       Pred. No. 10;
Mismatches
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Mismatches
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US-09-431-517-3422/c ; Sequence 3422, Application US/09431517

RESULT 12

Gaps

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100.0%; Score 20; DB 34; Length 449; 100.0%; Pred. No. 10; cive 0; Mismatches 0; Indels
CURRENT APPLICATION NUMBER: US/09/909,629
CURRENT FILING DATE: 2001-07-19
PRIOR APPLICATION NUMBER: 09/306,350
PRIOR FILING DATE: 1999-05-07
NUMBER OF SEQ ID NOS: 41304
SOFTWARE: DC CT_1 Version 1.1
SEQ ID NO 30669
LENGTH: 449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        completed: January 12, 2003, 12:04:06
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Matches 20; Conservative
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Matches 20; Conservative
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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Job time: 3167 secs
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APPLICANT: Dickson, Mark C.

APPLICANT: Dranac, Radoje T.

APPLICANT: Jones, Lee W.

APPLICANT: Stache-Crain, Birgit

TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained

TITLE OF INVENTION: From Various CDNA Libraries

FILE REFERENCE: 20411-776

CURRENT APPLICATION DIMBER: US/09/306,350A

CURRENT PILLING DATE: 1999-05-07

NUMBER OF SEQ ID NOS: 41304

SEQ ID NO 30669

LENGTH: 449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 100.0%; Score 20; DB 17; Length 449; Best Local Similarity 100.0%; Pred. No. 10; Matches 20; Conservative 0; Mismatches 0; Indels
                APPLICANT: Hyseq, inc

TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED

TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES

FILE REFERENCE: 20411-751.CON1

CURRENT APPLICATION NUMBER: US/09/431,517

CURRENT FILING DATE: 1999-11-01

EARLIER APPLICATION NUMBER: US 09/170,294

EARLIER FILING DATE: 1998-10-13

NUMBER: OF SEQ ID NOS: 31760

SOFTWARE: FASESEQ for Windows Version 3.0

SEQ ID NO 3422
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GENERAL INFORMATION:
APPLICANT: Dickson, Mark C.
APPLICANT: Drmanac, Radoje T.
APPLICANT: Labat, Ivan
APPLICANT: Stache-Crain, Birgit
TILLE OF INVENTION: Novel Nucleic Acid Sequences Obtained
TITLE OF INVENTION: From Various cDNA Libraries
FILE REFERENCE: 20411-776
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-306-350A-30669/c; Sequence 30669, Application US/09306350A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                   ; NAME/KEY: misc_feature
; LOCATION: (1)...(443)
; OTHER INFORMATION: n = A,T,C or G
US-09-431-517-3422
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             73 TTTCAACCAGATGGTCATTC 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 TTTCAACCAGATGGTCATTC 20
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US-09-306-350A-30669
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US-09-909-629-30669/c
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RESULT 15
US-09-824-557-3555/c
; Sequence 3555, Application US/09824557
; GENERAL INFORMATION:
; APPLICANT: Shyjan, Andrew W.
; TITLE OF INVENTION: THEREFOR;
; TITLE OF INVENTION: WOWER: US/09/824,557
; CURRENT FILING DATE: 2001-04-02
; PRIOR APPLICATION NUMBER: US 60/193,438
; PRIOR FILING DATE: 2000-03-31
; NUMBER OF SEQ ID NOS: 9773
; SEQ ID NO 3555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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100.0%; Pred. No. 10;
tive 0; Mismatches 0;
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OTHER INFORMATION: n = A,T,C or G
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APPLICANT: Henderson, Robert A.
APPLICANT: Wang, Tongtong
APPLICANT: Watanabe, Yoshihiro
APPLICANT: Kalos, Michael D.
APPLICANT: Sleath, Paul R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Matches 20, Conservative
; ORGANISM: Homo sapiens
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US-10-283-017-981
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               TYPE: DNA
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Sequence 20756, A
Sequence 20756, A
Sequence 20755, A
Sequence 20750, A
Sequence 20750, A
Sequence 20749, A
Sequence 20749, A
Sequence 20751, A
Sequence 20753, A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20758, A Sequence 20748, A Sequence 20748, A Sequence 92, Appl Sequence 20760, A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 981, App
                                                                                                   January 12, 2003, 09:51:30 ; Search time 168 Seconds (without alignments) 110.212 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /cgn2 6/ptodata/1/pna/DSO NEW COMB.seq:*

2: /cgn2_6/ptodata/1/pna/USO6_NEW COMB.seq:*

3: /cgn2_6/ptodata/1/pna/USO7_NEW COMB.seq:*

4: /cgn2_6/ptodata/1/pna/USO8_NEW COMB.seq:*

5: /cgn2_6/ptodata/1/pna/USO8_NEW COMB.seq:*

6: /cgn2_6/ptodata/1/pna/USO8_NEW COMB.seq:*

7: /cgn2_6/ptodata/1/pna/USO8_NEW COMB.seq:*
            GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-724-676A-20756
US-09-724-676A-20755
US-09-724-676A-20755
US-09-724-676A-20750
US-09-724-676A-20750
US-09-724-676A-20750
US-09-724-676A-20752
US-09-724-676A-20751
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US-09-724-676-20760
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-09-724-676A-20758
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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17917, Ap
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5041, Ap
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APPLICANT: Henderson, Robert A.
APPLICANT: Henderson, Robert A.
APPLICANT: Watanabe, Yoshihiro
APPLICANT: Kalos, Michael D.
APPLICANT: Sleath, Paul R.
APPLICANT: Sleath, Paul R.
APPLICANT: Retter, Marc W.
APPLICANT: Panger, Carrer, Darrick
APPLICANT: Fanger, Cary R.
APPLICANT: Bangur, Chaitanya S.
APPLICANT: Carrer, Darrick
APPLICANT: McNabb, Andria
APPLICANT: McNabb, Andria
TITLE OF INVENTION: CMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210.14.7803PC
CURRENT APPLICATION NUMBER: PCT/USO2/34777
CURRENT FILING DATE: 2002-10-28
NUMBER OF SEQ ID NOS: 2157
SOFTWARE FREESE FREESE
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US-09-724-676A-20760
US-10-203-138A-10274
US-10-203-138A-10274
US-10-196-793A-22
US-09-513-999C-18126
US-09-531-113-37959
US-09-531-113-37959
US-09-531-113-37959
US-09-531-113-37959
US-09-531-113-7917
US-09-724-676-5046
US-09-724-676-5046
US-09-724-676-5040
US-09-724-676-5040
US-09-724-676-5047
US-09-724-676-5049
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Watanabe, Yoshihiro
Kalos, Michael D.
Sleath, Paul R.
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Sequence 20756, Application US/09724676A
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative spl
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: PatentIn version 3.2
LENGTH: 1317
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; SEQ ID NO 20756
; LENGTH: 1317
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-724-676-20756
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; LENGTH: 550
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-283-017-981
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Best Local Similarity
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TYPE: DNA
ORGANISM: Homo sapiens
-09-724-676A-20756
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CURRENT APPLICATION NUMBER: US/10/283,017
CURRENT FILING DATE: 2002-10-28
NUMBER OF SEQ ID NOS: 2157
SOFTWARE: FastSEQ for Windows Version 4.0
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Bangur, Chaitanya S.
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Pred. No. 0.47;
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US-09-724-676-20750/c
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                                                                                                                                   Sequence 20750, Application US/09724676 GENERAL INFORMATION:
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TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97272
SOFTWARE: Patentin version 3.2
SEQ ID NO 20755
LENGTH. 1362
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: PatentIn version 3.2
SEQ ID NO 20750
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TITLE OF INVENTION: Variants of alternative
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 13
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TYPE: DNA
ORGANISM: Homo sapiens
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100.0%; Pred. No. 0.54;
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US-09-724-676A-20752/c
iSequence 2 0752, Application US/09724676A
; Sequence 2 0752, Application US/09724676A
; GENERAL INFORMATION:
    APPLICANT: Compugen LTD
; TITLE OF INVENTION: Variants of alternative splicing; FILE REFERENCE: 129181.4 Compugen
; CURRENT APPLICATION NUMBER: US/09/724,676A
; CURRENT FILING DATE: 2000-11-28
; NUMBER OF SEQ ID NOS: 97222
; SOFTWARE: PatentIn version 3.2
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Sequence 20752, Application US/09724676

GENERAL INFORMATION:

APPLICANT: Compugen LTD

TITLE OF INVENTION: Variants of alternative splicing

FILE REFERENCE: 129181.4 Compugen

CURRENT APPLICATION NUMBER: US/09/724,676

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 97222

SOFTWARE: Patentin version 3.2

SEQ ID NO 20752

LENGTH: 3610
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CURRENT APPLICATION NUMBER: US/09/724,676A CURRENT FILING DATE: 2000-11-28 NUMBER OF SEQ ID NOS: 97222 SSCYTWARE: Patentin version 3.2 SEQ ID NO 20749 LENGTH: 1836
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Pred. No.
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Matches 20; Conservative 0;
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Matches 20; Conservative
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Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                       TYPE: DNA
CORGANISM: Homo sapiens
US-09-724-676A-20749
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CORGANISM: Homo sapiens
US-09-724-676-20752
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; ORGANISM: Homo sapiens
US-09-724-676A-20752
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US-09-724-676-20751/c
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100.0%; Score 20; DB 5; Length 1791;
Best Local Similarity 100.0%; Pred. No. 0.56;
Matches 20; Conservative 0; Mismatches 0; Indels
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GENERAL INFORMATION:
FAPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
GURRENT APPLICATION NUMBER: US/09/724,676A
CUBRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patentin version 3.2
SEQ ID NO 20750
LENGTH: 1791
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Sequence 20749, Application US/09724676

GENERAL INFORMATION:

APPLICANT: Compugen LTD

TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen

CURRENT APPLICANION NUMBER: US/09/724,676

CURRENT FILING DATE: 2000-11-28

NUMBER OF SEQ ID NOS: 97222

SOFTWARE: Patentin version 3.2

SEQ ID NO 20749

LENGTH: 1836
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TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
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; Sequence 20749, Application US/09724676A
; GENERAL INFORMATION:
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Best Local Similarity 100.0
Matches 20; Conservative
                                          ; ORGANISM: Homo sapiens
US-09-724-676-20750
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US-09-724-676A-20750
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; ORGANISM: Homo sapiens
US-09-724-676-20749
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Best Local Similarity
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LENGTH: 1791
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Sequence 20751, Application US/09724676
GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patentin version 3.2
SEQ ID NO 20751
LENGTH: 3677
TYPE: DNA
ORGANISM: Homo sapiens
US-09-724-676-20751
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US-09-724-676-20754/c
; Sequence 20754, Application US/09724676
; Sequence 20754, Application US/09724676
; GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVERNION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
; CURRENT APPLICATION NUMBER: US/09/724,676
; CURRENT FILING DATE: 2000-11-28
; NUMBER OF SEQ ID NOS: 97222
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 20754
; LENGTH: 3697
; Type: DNA
; ORGANISM: Homo sapiens
US-09-724-676-20754
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US-09-724-676A-20751/c
; Sequence 20751, Application US/09724676A
; GENERAL INFORMATION:
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TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676A
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SOFTWARE: PatentIn version 3.2
SEQ ID NO 20751
LENGTH: 3677
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Matches 20; Conservative 0
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Search completed: January 12, 2003, 12:07:16 Job time : 170 secs

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January 12, 2003, 09:41:35 ; Search time 2677 Seconds (without alignments) 217.429 Million cell updates/sec
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Description	6330 Sequence	6663 human ST	68271 Sequenc	iomo sapie	 sap	Times fore	AL832961 Homo sapi	AX281683 Sequence	AC067931 Homo sapi	AJ132004 Rhizobium	AL591790 Sinorhizo	AC102801 Homo sapi	AC084336 Homo sapi	ACUITESZ HOMO SAPI	AC103694 Homo sani	AL713916 Mouse DNA	AC091767 Homo sapi	AL606668 Oryza sat	Mis		Mus mu	Rattus	ACUS4333 RACTUS NO AC111560 Rattus no	Homo sa	Rattus n	AL808136 Mus muscu	Rattus	Coriolus	Rattus n	eugenii	AA481431 Sequence AR007225 Semience	Sequenc	Sequenc	80	Sequence 1	אָ מ	ט כ			DNA linear PAT 16-MAY-2001						on kinase expression
DB ID	AR116	1 G06663	AX3682	HUMFAK	AKU9499			AX281683			SME59179					0	AC091767									2 AL808136	_						6 AR170011			7 7	AR069	ALIGNMENTS		20 bp	patent US 6133031.	:14096652			£ (1)	e inhibition of focal adhesion
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Primer B: AGTTTAGGAATTAAGATGTCL
STS size: 350
PCR Profile:
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                                                                                                    Prepared with primer pairs derived Location/Qualifiers
                                                                                                                                                                                                                                                                                        Protocol
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Homo sapiens STSs derived from sequences in dbEST and
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  117
                                                                                                                                                                                                                                              Template: 10 ng
Primer: each 5 pM
dNTPs: each 4 nM
                                                                                                                                                                   MgCl2: 1.5 mM
KCl: 50 mM
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Total Vol: 20 ul
                                                                                                                                                       Tris-HCL: 10 mM
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ilarity 100.0%;
Conservative (
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PCR Cycles: 35
Thermal Cycler:
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Annealing: 56 degrees C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                thudson@genome.wi.mit.edu
  ք
                         /organism="Homo sapiens"
/db_xref="taxon:9606"
/map="951_H_5; 805.4 cR fi
14. 363
14. 33
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complement(339..363)
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5 c 3 g
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RESULT 3
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                                                                                                                                               Andre, E. and Becker-Andre, M. Expression of an N-terminally adhesion kinase in brain
                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 3052)
                                                                                                                                                                                                                                                                                                                                                                                                                                               20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                        L05186
L05186.1 GI:182394
                                                                                                                                                                                                                                                                                                     HUMFAK 3052 bp Homo sapiens focal adhesion kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent: WO 0204514-A 981 17-JAN-2002; CORIXA CORPORATION (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                    Biochem. Biophys. Res. Commun.
                                                                                                                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                              focal adhesion kinase; protein-tyrosine
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                                                                                          Location/Qualifiers
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            /dev_stage="fetal"
15. _2654
                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
/tissue_type="brain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /db_xref="taxon:9606"
113 c 121 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           organism="Homo sapiens"
EC_number="2.7.1.1"
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                                                                                                                                                               truncated
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function="protein/tyrosine kinase"

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APGHLGSLASLSSPADSYNEGVKLKPQEISPPTANLDRSNDKVYENVTGLVKAVIEM
SSKIQPAPPEEYVPMVKEVGLALRTLLATVDETIPLLPASTHREIEMAQKLLNSDLGE
LINKMKLAQQYVMTSLQQEYKKQMLTAAHALAVDAKNLLDVIDQARLKMLGQTRRH"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ninomiya, K., Wagatsuma, M., Kanda, K., Kondo, H., Yokoi, T.,
Kodaira, H., Furuya, T., Takahashi, M., Kikkawa, E., Omura, Y., Abe, K.,
Kamihara, K., Katsuta, N., Sato, K., Tanikawa, M., Yamazaki, M.,
Sugiyama, T., Irie, R., Otsuki, T., Sato, H., Wakamatsu, A., Ishi, S.,
Yamamoto, J., Isono, Y., Kawai-Hio, Y., Sato, K., Nishikawa, T.,
Kimura, K., Yamashita, H., Matsuo, K., Nakamura, Y., Sekine, M.,
Kikuchi, H., Muzakawa, K., Kanehori, K., Takahashi-Fujii, A.,
Oshima, A., Sugiyama, A., Kawakami, B., Suzuki, Y., Sugano, S.,
Magahari, K., Masuho, Y., Magai, K. and Isogai, T.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Homo sapiens hippocampus CDNA to mRNA, clone_lib:BRHIP2
clone:BRHIP2012923.
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                                                                             tissue type="Brain, hippocampus"
/clone_lib="NIH_MGC_95"
                /db_xref="taxon:9606"
/clone="MGC:33483 IMAGE:4812915"
                                                                                                   /clone_lib="Nun_ruc_____/lab_host="DH10B"
/note="Vector: pBluescript"
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KEYWORDS
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/codon_start=1
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IPPLLARASTHREE BMAQKLLNSDLGELINKMKLAQQCYVMTSLQQEYKKQMLTRAALANV
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Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
Toshi Jibrary Preparation: Michael J. Brownstein (NHGRI) & Shiraki Cobh Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Institute for Systems Biology
DNA Sequencing by: Institute for Systems Biology
Contact: amadan@systemsbiology.org
Anup Madan, Jessica Fahey, Erin Helton, Mark Ketteman, Anuradha Madan, Stephanie Rodrigues, Amy Sanchez and Michelle Whiting
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
1 (bases 1 to 3199)
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Submitted (29-APR-2002) National Institutes of Health, Mammalian
Gene Collection (MGC), Cancer Genomics Office, National Cancer
Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Clone distribution: MGC clone distribution information can be for through the I.M.A.G.E. Consortium/LiNL at: http://image.llnl.gov Series: IRAK Plate: 46 Row: d Column: 12. This clone was selected for full length sequencing because it passed the following selection criteria: Hexamer frequency ORF
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Best Local Similarity 100.0
Matches 20, Conservative
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RESULT 5 BC028733/c

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COUNT

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ACCESSION

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ORGANISM

KEYWORDS SOURCE

AUTHORS TITLE JOURNAL

REMARK

COMMENT

REFERENCE

/organism="Homo sapiens"

source

FEATURES

PRI 15-JUL-2002 highly similar

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DEFINITION
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                            RESULT 8
HUMFAKX/c
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AUTHORS
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AR116314/c
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HUMFAKX 3791 bp mRNA
Human focal adhesion kinase (FAK) mRNA,
L13616
                                                                                                                                                                                                                                                                                                                                                                                                                               3791 bp
Sequence 1 from patent US 6133031.
AR116314
AR116314.1 GI:14096676
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Monia, B.P. and Gaarde, W.A.
Antisense inhibition of focal adhesion kinase
Patent: US 6133031-A 1 17-OCT-2000;
Location/Qualifiers
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NPLVSSNDCVKLGDFGLSRYMEDSTYYKASKGKLPIKMMAPESINFRRFTSASDVMMF 
GVCMWEILMHGVKPFQGVKNNDVIGRIENGERLPMPPNCPFLYSLMTKCWAYDPSRR 
RRFTELKAQLSTILEEEKAQQEERMRMESSRQATVSNDSGGSDEAPFKRESRPGYYBEP 
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                                                                                                                                                                                                                                                  /organism="unknown"
849 c 924 g
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EYKKQMLTAAHALAVDAKNLLDVIDQARLKMLGQTRPH"
a 697 c 741 g 837 t
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RESULT 9
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1 (bases 1 to 3791)
Whitney,G.S., Chan,P.Y., Blake,J., Cosand,W.L., Neubauer,M.G., Aruffo,A. and Kanner,S.B.
Aruffo,A. and Kanner,S.B.
Human T and B lymphocytes express a structurally conserved focal adhesion kinase, pp125FAK
DNA Cell Biol. 12 (9), 823-830 (1993)
                                                                                                                                                                                                                                                                            20;
Homo sapiens mRNA; cDNA
AL832961
AL832961.1 GI:21733549
                                                                     HSM804272
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Homo sapiens (tissue library: lambda ft
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DQVDOBIALKLGCLEIRRSYWEMRGNALEKKSNYEVLEKDVGLKRFFPKSLLDSVKAK
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n clone DKFZp66600110)
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                                                                                                                                                                                                         SOURCE
                                                                                                                                                                                                                                     German Genome Project.

This clone (DKFZp66600110) is available at the RZPD in Berlin.
Please contact the RZPD: Ressourcentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de Further
information about the clone and the sequencing project is available
at http://mips.gsf.de/proj/cDNA/.
Location/Qualifiers
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                                                   Mammalia; Eutheria; Primates; Catarrhin; Hominidae; Homo.
Mammalia; Eutheria; Primates; Catarrhin; Hominidae; Homo.
I (bases 1 to 4431)
Ansorge,W., Wirkner,U., Mewes,H.W., Weil,B. and Wiemann,S.
Direct Submission
Submitted (09-JUL-2002) 1, D-85764 Neuherberg, GERMANY
Clone from S. Wiemann, Molecular Genome Analysis; German Cancer Research Center (DKF2); Email s. wiemann@dkf2-heidelberg.de; sequenced by EMBL (European Molecular Biology Laboratories, Heidelberg/Germany) within the cDNA sequencing consortium of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                             /tissue_type="stomach"
/clone_lib="666 (synonym: hsto2). Vector pSport1; host
DH108; sites Not1 + Sal1"
/dev stage="adult"
4383. .4388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shiffman, D., Somogyi, R., Lawn, R., Seilhamer, J.J., Porter, G.J., Mikita, T. and Tai, J. Genes expressed in foam cell differentiation Patent: WO 017389-A 92 18-OCT-2001; Incyte Genomics, Inc. (US)
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974 c 1045 g 1136 t
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Sequence 92 from Patent WO0177389.
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/db_xref="taxon:9606"
/clone="DKFZp66600110"
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AXZ81683.1 GI:16608934
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                                                                                                                                                                                                                                                                                                  Submitted (24-FEB-1999) Sanjuan J., Microbiologia del Suelo y Sistemas Simbioticos, Estacion Experimental del Zaidin-CSIC, Profesor Albareda 1, E-18008 Granada, SPAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 3974)
Sanjuan-Pinilla,J.M., Olivares,J. and Sanjuan,J.
The Rhizobium meliloti leuA gene is essential for
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Sinorhizobium meliloti
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                                                                                                                                                                                                                                                                                                                                                            Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                 Sanjuan,
                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rhizobiaceae; Sinorhizobium.
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This sequence will be replaced by the finished sequence as soon as it is available and the accession number will be preserved.

1 206467: contig of 206467 bp in length.
Location/Qualifiers
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------ Project Information
Center project name: L10082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Center clone name: 642_A_1
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                /codon_start=1
/trans1_table=11
/product="CLP protease proteolytic
/protein_id="CAB39976.1"
/db_xref="GI:4539549"
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136. .723
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/clone lib="RPCI-11 Human Male BAC"
50155 c 48369 g 50320 t 4 4
translation="MRNDDDQEEKKTELPLGKETEANLFKSRSIFIYGTITQELAQKV"
                                                                                                                  gene="clpP"
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Sinorhizobium meliloti 1021 complete
AL591790 AL591688
AL591790.1 GI:L507527n
                             1 (bases 1 to 323450)
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                                                                                                                                                                                                                                                                                                                                                             Sinorhizobium meliloti.
Sinorhizobium meliloti
                                                                                                                                                                                                                                                                             Bacteria; Proteobacteria; alpha subdivision; Rhizobiaceae group; Rhizobiaceae; Sinorhizobium.
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Pred. No. 1.5e+02;
D; Mismatches 1;
sequence of
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                                     Ramsperger, U., and Galibert, F.
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Gene name confidence : hypothetical
predicted by Codon usage
predicted by Homology
predicted by FrameD"
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Gene name confidence : hypothetical
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Plantes-Microorganismes, UMR215-CNRS-INRA, BP27, F-31326 Castanet,
France, Laboratoire de Genetique et Developpement UMR6051-CNRS,
Facult de Medecine, 2 avenue du Pr. Leon Bernard, F-35043 Rennes,
France, GATC GmbH, Fritz-Arnold-str. 23, D-78467 Konstanz, Germany,
Universitet Bielefeld, Biologie IV (Genetik) Universiteatstr 25,
Universite Catholique de Louvain, Place Croix du Sud 2, Bte 20,
B-1348 Louvain-la-Neuve, Belgium, Unite de Microbiologie, Faculte
des Sciences Agronomiques de Gembloux, Avenue Marechal Juin 6,
B-5030 Gembloux, Belgium E-mail:Jerome Gouzy@toulouse.inra-fr
http://sequence.toulouse.inra-fr/meliloti.html
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                                                                                                                                                                                                                                           Direct Submission
Submitted (26-JUL-2001) Gouzy J., Submitted on behalf of the MELILO
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21396507
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1707. .2996
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/gene="SWc01576"
/function="small molecule metabolism"
/note="Product confidence : putative
Gene name confidence : putative
Gene name confidence : hypothetical.
predicted by Codon usage
predicted by Homology
predicted by FrameD"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 organism="Sinorhizobium meliloti"
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/evidence=not_experimental
/transl_table=11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (protein id="CAC46788.1"
| db xref="GI:15075232"
| db_xref="SPTREMBL:Q92NJ3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   predicted by Codon usage predicted by Homology predicted by FrameD"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /strain="1021"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /gene="SMc01575"
155. .1519
                                                                                                                                                                  (bases 1 to 323450)
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                                                                                                                                                                                                                                                                                                                                    EU Consortium
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                                                                                                                                11481430
                                                                                                                                                                                                                 Gouzy, J
                                                                                                                                                                                                        AUTHORS
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                                                                            MEDLINE
PUBMED
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VERSION
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TTCCAACCAGATGGTCATTC 183347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19;
Birren, B., Linton, L., Nusbaum, C., Lander, E., Ali, A., Allen, N., Anderson, S., Barna, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B., Brown, A., Camarata, J., Campopiano, A., Chang, J., Chazaro, B., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cook, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Faro, S., Ferreira, P., Fitzhugh, W., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Gord, S., Goyette, M., Graham, L., Grand-Pierre, N., Hagos, B., Heaford, A., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C., Kamat, A., Karatas, A., Kells, C., LaRocque, K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 119671)
Birren,B., Linton,L., Nusbaum,C. and Lander,E.
Homo sapiens chromosome 8, clone RP11-960P16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AC102801
AC102801.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens chromosome 8, clone RP11-960P16, complete sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AC102801
                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                      (bases 1 to 119671)
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/db
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/gene="SMC01594"
/function="miscellaneous; hypothetical/global homology"
/note="Product confidence : hypothetical
Gene name confidence : hypothetical
predicted by Codon usage
predicted by Homology
predicted by FrameD"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ILVGERMVRAFIEAGVPADVFQNLFLDHDTTAALIAAKSFDFINFTGSVEGGRSIERA AAGTFTGLGLELGGKDPGYVMEDADLDAAVDTLMBGATYNSGCCCGIERIYVHESLY DAFVEKSVAWUSNYKLGNPLDPETTLGPMANKFFAATVROVADAISKGARALIDPKLFPQDDGGAYLAPQVLVDVDHSWEFMREFTFGPAVGIMKVKSDAEAIELMNDSRYGINSKULGHRUTTKDAERAARIGRDLETGTVFMNRADYLDPALCWTGVKETGRGGSLSVLGFHNLTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        complement (8131. .9678)
/gene="SMc01597"
/function="cell processes; transport of small molecules;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  complement(8131./gene="SMC01597"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino acids, amines, peptides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KEVREIPHTLREAGEALSGSKMLRAAFGEEVVDHYVHAAEWEQQEYDRRVTDWEVARG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /product="CONSERVED HYPOTHETICAL PROTEIN"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             codon_start=1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GI:20377028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92.0%;
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Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            119671 bp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   _experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA
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                                                                            repeat_region
repeat_region
                                                                                                                                                     repeat_region
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RS Birren, B., Linton, L., Nusbaum, C., Lander, E., Ali, A., Allen, N., Barren, B., Linton, L., Nusbaum, C., Lander, E., Ali, A., Allen, N., Anderson, S., Barna, N., Bastien, V., Bloom, T., Boguslavkiy, L., Boukhgalter, B., Brown, A., Camarata, J., Campopiano, A., Chang, J., Cook, P., Dearellano, K., Dewar, K., Diaz, J.S., Dodge, S., Cook, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Ferreira, P., FitzHugh, W., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Gord, S., Goyette, M., Graham, L., Grand-Pierre, N., Hagos, B., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C., Kamat, A., Karatas, A., Kals, C., LaRocque, K., Landazares, R., Landers, T., Lehoczky, J., Levine, R., Lindblad-Toh, K., Liu, G., MacLean, C., Macdonald, P., Major, J., Marquis, N., Matthews, C., McCarthy, M., McEwan, P., McKernan, K., Meddrim, J., Meneus, L., Mihova, T., Mlenga, V., Murphy, T., Naylor, J., Nguyen, C., Nicol, R., Norbu, C., Norman, C.H., O'Connor, T., O'Donnell, P., O'Neil, D., Oliver, J., Peterson, K., Phunkhang, P., Pierre, N., Pollara, V., Raymond, C., Retta, R., Rieback, M., Riley, R., Rise, C., Rogov, P., Raymond, C., Retta, R., Rieback, M., Riley, R., Rise, C., Rogov, P., Roman, J., Rosetti, M., Roy, A., Santos, R., Schauer, S., Schupback, R., Seaman, S., Severy, P., Spencer, B., Stange-Thomann, N., Stojanovic, N., Strauss, N., Subramanian, A., Talamas, J., Tesfaye, S., Theodore, J., Viel, R., Vo, A., Wilson, B., M., X., Wyman, D., Ye, W.J., Young, G., Direct Submission
Direct Submission

At., Submitted (Ol. May, 2002) Whitehead Institute/MTT Center for Conven
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MacLean,C., Macdonald,P., Major,J., Marquis,N., Matthews,C.,
McCarthy,M., McEwan,P., McKernan,K., McDheeters,R., Meldrim,J.,
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Raymond,C., Retta,R., Rieback,M., Riley,R., Rise,C., Rogov,P.,
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Seaman,S., Severy,P., Spencer,B., Stange-Thomann,N., Stojanovic,N.,
Strauss,N., Subramanian,A., Talamas,J., Tesfaye,S., Theodore,J.,
Topham,K., Travers,M., Travis,N., Trigillo,J., Vassiliev,H.,
Viel,R., Vo,A., Wilson,B., Wu,X., Wyman,D., Ye,W.J., Young,G.,
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SP6 end overlaps project AC011632 (L2781) by 63.5 kilobases; overlaps project AC100782 (L19699) by 47.5 kilobases. We are submitting only the middle 119.7 kilobases of the project.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Research, 320 Charles Street, Cambridge, MA 02141, USA
on May 1, 2002 this sequence version replaced gi:17060911.
All repeats were identified using RepeatMasker:
Smit, A.F.A. & Green, P. (1996-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (23-NOV-2001) Whitehead Institute/MIT Center Research, 320 Charles Street, Cambridge, MA 02141, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (01-MAY-2002) Whitehead Institute/MIT Center for Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /rpt_family="L2"
405. .447
                                                                                /rpt_family="AluY"
295. .366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                             map="8"
                                                                                                                                                                                                                                                                                                                                                      organism="Homo sapiens"
|db_xref="taxon:9606"
                                                                                                                                                                                         clone_lib="RPCI-11
                                                                                                                                                                                                                                                                                                                      chromosome="8"
                                                                                                                                                                                                                                             clone="RP11-960P16"
                                                                                                                                                                                                 Human Male BAC"
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rpt_family="AT_rich"

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Complement (8168. .8342)

/ TPL family="HAL1"

8648. .8676

/ TPL family="HAL1"

Complement (8760. .9400)

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Complement (9662. .9939)

/ TPL family="HAL1"

Complement (10387. .10511)

/ TPL family="HAL1"

Complement (10387. .10511)
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complement(4609. .4939)
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4957. .5281
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Streen, B., Linton, L., Nusbaum, C., Lander, E., Abraham, H., Allen, N., Anderson, S., Barna, N., Bastien, V., Beda, F., Boguslavkiy, L., Boukhgalter, B., Erom, A., Burkett, G., Campoplano, A., Casile, A., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodges, S., Ferreira, P., FitzHugh, W., Gage, D., Galagn, J., Gardyna, S., Ginde, S., Goyette, M., Ilev, I., Johnson, R., Jones, C., Kann, L., Karates, R., Larkogue, K., Lamacares, R., Landers, T., Lehcozky, J., Levine, R., Lieu, C., Liu, G., Macdonald, P., Marquis, N., McCarthy, M., McEwan, P., McKernan, K., Morrow, J., Murphy, T., Naylor, J., Norman, C.H., O'Connor, T., O'Donnell, P., O'Neil, D., Olivar, T. M., Oliver, J., Peterson, K., Peirer, N., Piana, J., Roy, A., Santos, R., Schuuer, S., Severy, P., Sougnez, C., Spencer, B., Stange-Thomann, N., Stojanovic, N.,
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Homo sapiens chromosome 8, clone RP11-625F15, complete sequence.
AC084336
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1 (bases 1 to 156235)

Birran, B., Linton, L., Nusbaum, C. and Lander, E.

Homo sapiens chromosome 8, clone RP11-625F15
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http://ftp.genome.washington.edu/RM/RepeatMasker.html	Anderson, S., Barna, N., Bastlen, V., Boguslawiy, L., Boukingaller, B., Brown, A., Camarata, J., Campopiano, A., Chang, J., Chazaro, B., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cook, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Faro, S., Gord, S., Gord, S., Gayette, M., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Gord, S., Goyette, M., Graham, L., Grand-Pierre, N., Hagos, B., Heaford, A., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C., Kamat, A., Karatas, A., Kells, C., LaRocque, K., Jones, C., Manders, T., Lehoczky, J., Levine, R., Liu, G., MacLean, C., Macdonald, P., Major, J., Marquis, N., Matthews, C., McCarthy, M., McEwan, P., McKernan, K., McPheeters, R., Lauders, C., McCarthy, M., McEwan, P., McKernan, K., McPheeters, R., Meldrin, J., McCarthy, M., McEwan, P., McKernan, K., McPheeters, R., Meldrin, J., McCarthy, M., McEwan, P., McRernan, K., McPheeters, R., Meldrin, J., McCarthy, M., McEwan, P., McRernan, K., McPheeters, R., Meldrin, J., McCarthy, M., McBwan, P., McRerre, N., Pollara, V., Oliver, J., Peterson, K., Phinkhang, P., Pierre, N., Pollara, V., Raymond, C., Retta, R., Rieback, M., Riley, R., Rise, C., Rogov, P., Raymond, C., Rosetti, M., Roy, A., Santos, R., Schauer, S., Schupback, R., Seaman, S., Severy, P., Spencer, B., Stange-Thomann, N., Stojanovic, N., Strauss, N., Trayers, M., Talamas, J., Teffaye, S., Theodore, J., Viel, R., Vo, A., Wilson, B., Wu, X., Wyman, D., Ye, W. J., Young, G., Zainoun, J., Zembek, L., Zimmer, A. and Zody, M. Direct Submission Submitted (11-DEC-2001) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, M., 02141, USA On Dec 11, 2001 this sequence version replaced gi:15042917. All repeats were identified using RepeatMasker:	marata, J., Campopiano, A., Chang, J., Cha Colangelo, M., Collins, S., Collymore, A., Arellano, K., Dewar, K., Diaz, J.S., Dodge FitzHugh, W., Gage, D., Galagan, J., Gard rd, S., Goyette, M., Graham, L., Grand-Pie aford, A., Horton, L., Hulme, W., Iliev, I., mar, A., Karatas, A., Kells, C., LaRocque, , Landers, T., Lehoczky, J., Levine, R., I. Macdonald, P., Major, J., Marquis, N., Macdonald, P., McKernan, K., McPheeters, R., Ihova, T., Mlenga, V., Murphy, T., Naylor, rman, C.H., O'Connor, T., O'Donnell, P., O' eterson, K., Phunkhang, P., Pierre, N., Po eterson, K., Phunkhang, P., Pierre, N., Po eterson, K., Phunkhang, P., Pierre, N., Po eterson, K., Phunkhang, P., Fierre, N., Po eterson, K., Phunkhang, P., Schauer, S., setti, M., Roy, A., Santos, R., Schauer, S., setti, M., Roy, A., Santos, R., Schauer, S., setti, M., Roy, A., Santos, R., Schauer, S., very, P., Sougnez, C., Spencer, B., Stang Viel, R., Vo, A., Wilson, B., Wu, X., Wym inoun, J., Zembek, L., Zimmer, A. and Zody Johans, Street, Cambridge, MA 02141, to 15635) inton, L., Nusbaum, C., Lander, E., Ali, A., "Inton, L., Nusbaum, C., Lander, E., Ali, A.,	Strauss, N., Subramanian, A., Talamas, J., Tetraye, S., Theodore, J., Tirrell, A., Travers, M., Trigilio, J., Vassiliev, H., Viel, R., Vo, A., Wilson, B., Wu, X., Wyman, D., Ye, W.J., Young, G., Zainoun, J., Zimmer, A. and Zody, M. Direct Submission Direct Submission Submitted (25-0CT-2000) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA 3 (bases 1 to 156235) Birren, B., Linton, L., Nusbaum, C., Lander, E., Ali, A., Allen, N., Anderson, S., Barna, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B., Anderson, S., Barna, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B.,
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Search completed: January 12, 2003, 10:38:36 Job time : 2817 secs

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AAI43434
AAQ34687
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AAC65550 standard; DNA;
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            GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                          Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                    2185239 seqs, 1125999159 residues
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                                                                                                                                                                                                                                                                                                       Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                           - nucleic search, using sw model
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AAS15398
ABK38943
AAC65534
AAC5534
AAS15382
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                                                                                   2003, 09:40:15
                                                                                                                                                          20
                                                                                                                                                                                  IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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Query Match
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15398
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; focal adhesion kinase; FAK; melanoma; tumour metastasis; cancer; angiogenic disorder; retinal neovascularisation; cytost ophthalmological; antisense; phosphorothioate; ss.
                                                        treatment
                                                                           Antisense
                                                                                                              WPI; 2002-010103/01
                                                                                                                                                   Monia
                                                                                                                                                                                                                                                                                                                         09-JAN-2001; 2001US-0757100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human focal adhesion kinase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. Th antisense sequences, including the one shown here, can be used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        neovascularization, or for diagnosing and treating e.g. colon cancer
                                                                                                                                                                                        (MONI/) MONIA B P.
(GAAR/) GAARDE W A.
(NERO/) NERO P S.
                                                                                                                                                                                                                                                                19-AUG-1999; 99US-0377310
13-JUL-2000; 2000WO-US18999
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                                 suppression of Focal Adhesion Kinase expression for the of cancers of the breast, colon, mouth or skin (especially, and angiogenic disorders e.g. retinal neovascularisation
                                                                                                                                                   Gaarde WA,
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ilarity 100.0%;
Conservative 0
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/mod_b;
 19pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                    optionally bases 2'-methoxyethoxy 2'-MOE cytosines
                                                                                                                                                     Nero
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Pred. No. 0.78;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      also 5'methylcytosines"
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present invention relates to novel antisense compounds which can be d for modulating the expression of human focal adhesion kinase (FAK)

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RESULT 3
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Best Local
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29-AUG-2000;
08-SEP-2000;
26-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The antisense compounds comprise antisense oligonucleotides (8-30 nucleotides in length) targeted to the 5'-untranslated region, translational termination region or 3' untranslated region of a nucleic acid molecule encoding FAK. The antisense oligonucleotides can be used to inhibit the expression of FAK mRNA. The antisense oligonucleotides represent potential chemotherapeutic agents in the treatment of melanoma and the prevention of tumour metastasis. The antisense compounds and methods are useful for treating diseases associated with overexpression or constitutive activation of FAK. Such diseases include cancers of the breast, colon, mouth or skin (especially a melanoma), and anglogenic disorders such as retinal neovascularisation. AASI5383-AASI5422
The invention describes an isolated polynucleotide and polypeptide useful for stimulating and/or expanding T cells specific for a tumour protein for determining the presence of a cancer in a patient. A composition containing the polynucleotide and/or polypeptide is useful for treating a lung cancer in a patient. The polypeptide is useful for memoving tumour cells from a biological sample. The polynucleotide is
                                                                                                                             Novel polynucleotide encoding a lung stimulating and/or expanding T cells
                                                                                                                                                                         WPI; 2002-164634/21.
                                                                                                                                                                                                                              Wang T,
                                                                                                                                                                                                                                                                                                                                 06-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cDNA encoding lung tumour protein clone R0127:G07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABK38943;
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                                                                                                    Claim 1;
                                                                                                                                                                                                                  Marnerakis M,
                                                                                                                                                                                                                                                                                         03-MAY-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              represent the FAK antisense oligonucleotides of the invention
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                                                                                                                                                                                                  Watanabe Y, Henderson RA, Jo
is M, Carter D, Fanger GR, V
, Wang A, Fanger N, Switzer
                                                                                                  SEQ ID No 981; 223pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard; cDNA; 550
                                                                                                                                                                                                                                                                                      2000US-0677419;
2000US-0702705;
2000US-0736457;
2000US-0736457;
2000US-0849626;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene;
                                                                                                                                                                                                                                                                                                                                             2000US-0614124.
2000US-0651563.
2000US-0658824.
2000US-0671325.
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Pred. No.
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Vedvick TS,
zer A, Mcneill
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 response stimulator;
                                                                                                                             specific for a tumour protein
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ll PD, Clapper i
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(first entry)

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Antisense suppression of Focal Adhesion Kinase expression for the treatment of cancers of the breast, colon, mouth or skin (especially a melanoma), and anglogenic disorders e.g. retinal neovascularisation -
                                                                                                                                         Human; focal adhesion kinase; FAK; melanoma; tumour metastasis; cancer; angiogenic disorder; retinal neovascularisation; cytostatic; ophthalmological; ss.
                                                                                                               Human focal adhesion kinase (FAK) cDNA seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page -; 19pp; English.
                                AAS15382 standard; cDNA; 3791 BP.
                                                                                                                                                                                                                                                                                                           19-AUG-1999; 99US-0377310.
13-JUL-2000; 2000WO-US18999.
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                                                                                                                                                                                                                                                                                                                                                                                                           Gaarde WA,
                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-010103/01.
                                                                                                                                                                                                                                                                                                                                                  (MONI/) MONIA B P.
(GAAR/) GAARDE W A.
(NERO/) NERO P S.
                                                                                                                                                                                                                           US2001034329-A1.
                                                                                                                                                                                                 Homo sapiens,
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                                                                                     29-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                           Monia BP,
                                                          AAS15382;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. The antisense sequences can be used in the treatment of all of these.
also useful as probe or primer to detect the level of mRNA encoding a tumour protein. This sequence encodes a lung tumour associated protein or protein fragment, described in the method of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly
                                                                                                                                                                                                                                                                                                                                                                                                     Human, focal adhesion kinase, FAK, signal transduction, cancer, embryonic development disorder, angiogenic disorder, wound healing; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a number of phosphorothioate antisense
                                                                                                                                                                      Gaps
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Pred. No. 1.9;
                                                                                                                                      100.0%; Score 20; DB 24; Length 550; 100.0%; Pred. No. 1.4; ive 0; Mismatches 0; Indels
                                                                                                                                                                   0; Indels
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                                                                                                           Sequence 550 BP; 155 A; 113 C; 121 G; 161 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                             Human focal adhesion kinase coding sequence.
                                                                                   ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Column 31-40; 30pp; English.
                                                                                                                                                                                                                                                                                           AAC65534 standard; DNA; 3791 BP
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Best Local Similarity 100.
Matches 20. Conservative
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Matches 20; Conservative
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                                                                      rom WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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AAC65534/c
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The present invention relates to novel antisense compounds which can be used for modulating the expression of human focal adhesion kinase (FAK). The antisense compounds comprise antisense oligonuclectides (8-30 nucleotides in length) targeted to the 5'-untranslated region, translational termination region or 3' untranslated region of a nucleic acid molecule encoding FAK. The antisense oligonucleotides can be used to inhibit the expression of FAK mRNA. The antisense oligonucleotides represent potential chemcherapeutic agents in the treatment of melanoma and the prevention of tumour metaetasis. The antisense compounds and methods are useful for treating diseases associated with overexpression
                                                                                                                                                                                                                                                                                                                                                                                 breast, colon, mouth or skin (especially a melanoma), and angiogenic disorders such as retinal neovascularisation. The present cDNA sequence for human FAK is used to design the FAK antisense oligonucleotides
                                                                                                                                                                                                                                                                                                                                                     or constitutive activation of FAK. Such diseases include cancers of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Note: This sequence (Genbank accession number L13616) is not given in the patent but is acquired by the indexer from Genbank from the information provided in the patent.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 20; DB 24; Length 3791; 100.0%; Pred. No. 1.9;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 3791 BP; 1110 A; 849 C; 924 G; 908 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Fix
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of the invention.
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ID ABV2919
XX
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Best Local (
                                                                                                                                                                                                       Matches
                                                                                                                                                     3536
                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an isolated nucleic acid molecule (I) com a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate
           Human DNA sequence #92 expressed during
                                    14-FEB-2002
                                                                                       AAS94837
                                                                                                                                                                                                                                                     Sequence 4052 BP; 1199 A; 906 C; 971 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usef
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-662795/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-FEB-2000; 2000US-183319P.
16-MAR-2000; 2000US-189862P.
25-MAY-2000; 2000US-207454P.
09-JUN-2000; 2000US-211314P.
18-JUL-2000; 2000US-219007P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; prostate cancer; cytostati pharmacogenomic marker; gene; ss.
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                                                                                                                                                                                                                                                                                                                                                                                  cancer
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                                                                                                                                                                                                                                                                                                                               (E) (E)
                                                                                                                                                                                                                                                                                                                                                                   (d) assessing the efficacy of a therapy for inhibiting prostate cancer
                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                                                                                                   selecting a composition for inhibiting prostate cancer in a patient; assessing the prostate cell carcinogenic potential of a compound; determining whether prostate cancer has metastasized in a patient; assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            detecting
                                                                                                                                                                 TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sapiens
                                                                                                                                                     TTTCAACCAGATGGTCATTC 3517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prostate
                                                                                                                                                                                                                                                                              also useful as a pharmacodyanamic or pharmacogenomic marker
                                                                                                                                                                                                                                                                                                                                                                                  in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                      standard;
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                                                                                                                                                                                                                                                                                                                                                                                patient,
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                                    (first
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                                                                                      DNA;
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                                      entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; carcinogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monahan JE;
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Pred. No. 1.9;
                                                                                                                                                                                                       Mismatches
             foam cell differentiation
                                                                                                                                                                                                                                                      970 T; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            stage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of prostate
                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequences that are differentially expressed during foam cell differentiation. The polynucleotide sequences of the invention or a composition comprising these polynucleotides are useful as a high throughput method for detecting altered expression of one or more polynucleotides in a sample. The polynucleotides can be used in the diagnosis of disorders associated with foam cell development such as atherosclerosis, cerebral stroke, and cardiovascular disorders such as atherosclerosis, cerebral stroke, and cardiovascular disorders such as coronary artery disease. The polynucleotide sequences can also be used as PCR primers and probes. The polynucleotides of the invention are also useful in gene therapy. ARS99746-ARS9501 represent the human polynucleotide sequences of the invention which are differentially expressed during foam cell differentiation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3558
                                                                              02-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                  Human neuroblastoma expressed polynucleotide SEQ ID NO 3903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAI97828 standard; cDNA; 707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a sequences that are differentially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 145-146; 315pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Composition useful for diagnosis of conditions, disorders or diseases associated with atherosclerosis, comprises several polynucleotides that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; foam cell differentiation; atherosclerosis; cerebral stroke; cardiovascular disorder; coronary artery disease; gene therapy; ds.
                                                                                                                                                                                        WO200166719-A1
                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                       13-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-APR-2001; 2001WO-US11128
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                               neuroblastoma;
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                                                                              2001WO-JP01629
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                                                                                                                                                                                                                                                                                            malignancy; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 relates to the isolation of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВP
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in foam cell
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                                                                                                                                                                                                                                                                                            tumour marker; N-myc;
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07-MAR-2000; 2000JP-0159195

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Penn SG, Hanzel DK,
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                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                   WO200157271-A2.
                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                        01-FEB-2002
                                                                                                                                                                                                                                                                                                                                                        09-AUG-2001
                                                                                                                                                                                                                     ABA51579;
                                                                                  Query Match
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                                                                                                                                          The invention relates to novel genes (AAI93926-AAI97963) expressed in human neuroblastoma. The nucleic acids are applicable as a probe or primer in diagnosing the prognosis of human neuroblastoma, malignancy and susceptibility indicators or tumour markers for anti-cancer agents. The gene information for diagnosing prognosis is related to factors similar to that for N-myc and TrkA genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence encodes an aldo/ketoreductase (I) isolated from the white rot basidiomycete, Coriolus versicolor. The present invention also describes a method for detecting at least one sulphur-containing
                                                                     Nucleic acids originating in gene expressed in human neuroblastoma, useful as probe or primer in diagnosing prognosis of human neuroblastoma, malignancy and susceptibility indicator or tumour marker
                                                                                                                                                                                                                                                                                                                                                                                                                                      Aldo/ketoreductase; white-rot basidiomycete; methylbenzothiophene; AAD; aryl-alcohol dehydrogenase; enzyme; gene; ds.
                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                 Coriolus versicolor aldo/ketoreductase encoding DNA SEQ ID NO:1.
                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                      92.0%; Score 18.4; DB 22; Length 707; 95.0%; Pred. No. 9.8;
                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                   Sequence 707 BP; 200 A; 129 C; 149 G; 224 T; 5 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /product= "aldo/ketoreductase"
                                                                                                                                                                                                                                                Pred. No. 9.8;
                                                                                                                         Claim 1; Page 2870-2871; 2979pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A gene encoding a new protein and its use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
354..1229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 7-8; 12pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                       ABA93211 standard; DNA; 1311 BP
(CHIB-) CHIBA PREFECTURE, (HISM ) HISAMITSU PHARM CO LTD.
                                                                                                                                                                                                                                                                                                  431 Trrcaaccaggregreatre 450
                                                                                                                                                                                                                                                                                1 TTTCAACCAGATGGTCATTC 20
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                                                                                                                                                                                                                                                                                                                                                                                              17-APR-2002 (first entry)
                                                                                                                                                                                                                                    Query Match
Best Local Similarity 95.0
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a
                                                                                                     for anti-cancer agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Coriolus versicolor.
                                                   WPI; 2001-565584/63.
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(KUBI ) KUBOTA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-158675/21
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                               Nakagawara A;
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The invention relates to a spatially-addressable set of single exon nucleic acid probes for measuring gene expression in a sample derived from human breast and Br 47 cells. The method involves conteacting the probes with a collection of detectably labelled nucleic acids derived from mRNA of human breast, and then measuring the label bound to each probe of the microarray. The probes are useful for verifying the expression of regions of genomic DNA predicted to encode proteins. They are useful for gene discovery, and for encode proteins. They are useful for gene discovery, and for determining predisposition and/or prognosing breast disease. Gene expression analysis is useful for assessing the toxicity of chemical agents on cells. The microarray of this invention presents a far greater diversity of probes for measuring gene expression, with far less bias than expressed sequence tag microarrays. The method is suitable for rapid production of functional information from genomic sequence. The present sequence is a single exon nucleic acid probe of the invention.
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hydrocarbon compound selected from methylbenzothiophenes in which, in a process of assaying a compound by the expression of [1], the increase in the expression of [1] shows that the compound is the objective compound [1] and the gene encoding [1] can be used for decomposing noxious
                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                              compounds
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0
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                                                                                                                                                                                                                                                                                                                 DB 24; Length 1311;
58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; SEQ ID NO 10274; 327pp + sequence listing; English
                                                                                                                                                              substances, particularly sulphur-containing heterocyclic
                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human breast cell single exon nucleic acid probe #10274.
                                                                                                                                                                                                                                            Sequence 1311 BP; 340 A; 332 C; 391 G; 248 T; 0 other;
                                                                                                                                                                                                                                                                                                                                         100.0%; Preα. ...
've 0; Mismatches
                                                                                                                                                                                                                                                                                                                     Score 17;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABA51579 standard; DNA; 244 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-FEB-2000, 2000US-0180312.
26-MAY-2000; 2000US-0204456.
30-UTM-2000; 2000US-060840B.
03-AUG-2000; 2000US-023366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0234687.
04-OCT-2000; 2000GB-0234687.
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                                                                                                                                                                                                                                                                                                                     85.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                      17; Conservative
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RESULT 11
ABA36555
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                        measuring human gene expression in a sample derived from human heart. present sequence is one such probe. The probes may be used for predicting, measuring and displaying gene expression in samples derive from the human heart via microarrays. By measuring gene expression, the probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular systems. Cardiovascular disease, hypertension, cardiac arrhythmias and concentral heart disease, hypertension, cardiac arrhythmias and
                                                                                                                                                                                                                                                                                                                      Penn
                                     Sequence
                                                                      specification,
                                                                        Note: The sequence data for this patent did not form specification, but was obtained in electronic format
                                                                                                congenital heart disease
                                                                                                                                                                                                     The present invention relates to single exon nucleic acid probes
                                                                                                                                                                                                                              Claim 4;
                                                                                                                                                                                                                                                                  Single exon nucleic acid probes
                                                                                                                                                                                                                                                                                              WPI; 2001-488899/53
                                                                                                                                                                                                                                                                                                                                                                        04-OCT-2000;
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27-SEP-2000;
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26-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           congenital heart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; gene expression; heart; microarray; vascular syste cardiovascular disease; hypertension; cardiac arrhythmia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Probe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Note: The sequence data for this patent did not form part of printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                               (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                                         30-JUN-2000;
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                                                        ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                      SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TTTCAACCAGATGTACATTC 137
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                                                                                                                                                                                                                                                                                                                                               MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                             SEQ ID No 15021; 530pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                                      Hanzel
                                                                                                                                                                                                                                                                                                                                                                     ; 2000US-0180312
2000US-0207456
; 2000US-0608408
; 2000US-0632366
; 2000US-0234369
; 2000US-0234369
; 2000US-0234369
; 2000US-0236359
; 2000US-0024263
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                     50
                                                                                                                                                                                                                                                                                                                      DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene expression analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                     A; 59 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    84.0%;
90.0%;
84.0%;
90.0%;
                                                                                                                                                                                                                                                                                                                       Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      heart; microarray; vascular system; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          244
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Score
Pred.
                                     36 G;
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Pred. No. 5
                                                                                                                                                                                                                                                                                                                       Rank DR;
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                                                                                                                                                                                                                                                                   for analyzing gene expression in human
16.8;
No. 56;
                                     99 T; 0 other;
           DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              in human heart
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             22;
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           Length
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                                                                          part of the printed directly from WIPO
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              244;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cell sample.
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                                                                                                                                                                 derived
                                                                                                                           system
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RESULT 13
ABA46482
ID ABA46
XX
AC ABA46
XX
DT 01-FE
XX

ABA46482

standard;

DNA; 400 BP

01-FEB-2002 ABA46482;

(first entry)

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RESULT 12
AAK43716
ID AAK43
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                                           Matches
                                                     Best Local
                                                              Query Match
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                                                                                                       The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cancer such as lymphoma, leukaemia and myeloma. The present sequence is one of the probes of the invention.
                                                                                    Sequence
                                                                                                                                                                                 Example 4; SEQ ID NO:
                                                                                                                                                                                                                                                                                                                        03-AUG-2000;
21-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                       04-FEB-2000;
26-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                            09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; bone marrow expressed exon; gene expression analysis; probe;
microarray; cancer; leukaemia; lymphoma; myeloma; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-NOV-2001
                                                                                                                                                                                                                                                          Penn
                                                                                                                                                                                                                                                                                                                                             30-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human bone marrow expressed single exon probe SEQ ID NO:
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                                                                                                                                                                                                                                                                                (MOLE-)
 118
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                                                                                                                                                                                                                                                          SG,
                     TTTCAACCAGATGGTCATTC 20
TTTCAACCAGATGTACATTC 137
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                                                                                                                                                                                                     genome-derived single exon nucleic acid probes useful for ting gene expression in human bone marrow -
                                          l Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18;
                                                                                                                                                                                                                                                                                MOLECULAR DYNAMICS INC
                                                                                    244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                          Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                        2000US-0608408.
2000US-0632366.
2000US-0234687.
                                           Conservative
                                                                                    BP;
                                                                                                                                                                                                                                                                                                                                                                                       2001WO-US00668
                                                                                                                                                                                                                                                                                                    2000US-0236359.
2000GB-0024263.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                   50 A; 59 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                   84.0%;
90.0%;
                                                                                                                                                                                 18273; 658pp + Sequence Listing;
                                                                                                                                                                                                                                                          Chen W, Rank DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         244 BP
                                          0
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                                                    Score 16.8;
Pred. No. 5
                                                                                    36 G;
                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                    99 T;
                                                      56
                                                                                    0 other;
                                                               DB
                                                               22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2,
                                          Indels
                                                              Length
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                                                                                                                             treatment of cancers
                                                                                                                                                                                 English
                                                               244;
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                                          Gaps
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The present invention relates to single exon nucleic acid probes for measuring human gene expression in a sample derived from human heart. The present sequence is one such probes may be used for predicting, measuring and displaying gene expression in samples derived from the human heart via microarrays. By measuring gene expression, the probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular system e.g. cardiovascular disease, hypertension, cardiac arrhythmias and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Single exon nucleic acid probes for analyzing gene expression in human hearts -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; bone marrow expressed exon; gene expression analysis; probe; microarray; cancer; leukaemia; lymphoma; myeloma; ss.
     Human; gene expression; heart; microarray; vascular system; probe; cardiovascular disease; hypertension; cardiac arrhythmia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human bone marrow expressed single exon probe SEQ ID NO: 5265.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         84.0%; Score 16.8; DB 22; Length 400; 90.0%; Pred. No. 61;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 400 BP; 94 A; 97 C; 84 G; 125 T; 0 other;
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID No 5154; 530pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                             Rank DR;
                                                                                                                                                                                                                                                                                                                                                                         (MOLE-) MOLECULAR DYNAMICS INC
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                                                                                                                                                                                                                    04-FEB-2000; 2000US-0180312.
26-MAY-2000; 2000US-0207456.
30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0234687.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          344 TITCAACCAGATGTACAITC 363
                                         congenital heart disease; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 TTTCAACCAGATGGTCATTC 20
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Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-488899/53.
                                                                                                                WO200157274-A2
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                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              from human breast and BT 474 cells. The method involves contacting the probes for measuring gene expression in a sample derived from human breast and BT 474 cells. The method involves contacting the probes with a collection of detecrably labelled nucleic acids derived from mRNA of human breast, and then measuring the label bound to each probe of the microarray. The probes are useful for verifying the expression of regions of genomic DNA predicted to determining predisposition and/or prognosing breast disease. Gene expression analysis is useful for assessing the toxicity of chemical expression analysis is useful for assessing the toxicity of chemical expression analysis is useful for assessing the toxicity of chemical diversity of probes for measuring gene expression, with far less bias than expressed sequence tag microarrays. The method is suitable for caping production of functional information from genomic sequence. The present sequence data for this patent did not form part of the invention. Note: The sequence data for this patent did not form part of the contactly prince of the present sequence data for this patent did not form part of the contactly production of the present sequence data for this patent did not form part of the contactly production.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a spatially-addressable set of single exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New spatially-addressable set of single exon nucleic acid probes, useful for measuring gene expression in sample derived from human breast, comprises number of single exon nucleic acid probes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
0
                                      Human; microarray; single exon probe; gene expression; breast;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 84.0%; Score 16.8; DB 22; Length 400; Best Local Similarity 90.0%; Pred. No. 61; Matches 18; Conservative 0; Mismatches 2; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 5177; 327pp + sequence listing; English.
Human breast cell single exon nucleic acid probe #5177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                            Chen W, Rank DR
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                                                                                                                                                                                                                                                                                                                                                                                        (MOLE-) MOLECULAR DYNAMICS INC.
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2000US-0234687.
2000US-0236359.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-JAN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                            Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-496933/54
                                                          disease; cancer; ss
                                                                                                                             WO200157271-A2
                                                                                                                                                                                                                                                                                             03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
                                                                                             Homo sapiens
                                                                                                                                                                                                                                       04-FEB-2000;
26-MAY-2000;
                                                                                                                                                                                                                                                                            30-JUN-2000;
                                                                                                                                                               09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                          Penn SG,
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AC ABA26
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Search completed: January 12, 2003, 09:51:21 Job time : 255 secs
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Best Local Similarity 90.0
Matches 18; Conservative
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26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                                                                                                                                                                               The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cancers such as lymphoma, leukaemia and myeloma. The present sequence is one of the probes of the invention.
                                                                                                                                                      Sequence 400 BP; 94 A; 97 C; 84 G; 125 T; 0 other;
                                                                                                                                                                                                                                                                                                  Penn
                                                                                                                                                                                                                                                                         Example 4; SEQ ID NO: 5265; 658pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                           WPI; 2001-488900/53.
                                                                                                                                                                                                                                                                                                                                                                                             (MOLE-) MOLECULAR DYNAMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JAN-2001; 2001WO-US00668
                                                    344 TTTCAACCAGATGTACATTC 363
                                                                   1 TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                   SG,
                                                                                                                                                                                                                                                                                                                                                                   Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                     ; 2000US-0180312.
; 2000US-0207456.
; 2000US-0608498.
; 2000US-0632366.
; 2000US-0234687.
; 2000US-0236559.
; 2000US-0236559.
                                                                                                                  84.0%; Score 16.8; DB 22; Length 400; 90.0%; Pred. No. 61;
                                                                                                                                                                                                                                                                                                                                                                   Chen W,
                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                      Rank DR;
                                                                                                         Mismatches
                                                                                                        2;
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Appli Appli

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US-09-377-310-18

Sequence 18, Application US/09377310B

Patent No. 613301

GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Expression
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1, Application US/09377310B
Sequence 1, Application US/09377310B
Setent No. 6133031
GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT APPLICATION NUMBER: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO! SEQ ID NOT. 2.0
                                                     Sequence 2
Sequence 2
Sequence 2
Sequence 2
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Sequence 9
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Sequence 1
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US-08-664-596B-4
US-08-721-798A-1
US-08-961-527-75
US-08-724-394A-20
US-08-724-394A-21
US-08-724-394A-21
US-09-233-086-53
US-09-641-638-91
US-09-641-638-91
US-08-467-948A-1
US-08-467-948A-1
US-08-467-948A-1
US-08-887-798-1
US-09-178-252-20
US-09-357-071-39
US-09-357-071-39
US-08-263-346-1
US-09-357-071-39
US-08-263-346-1
US-09-357-071-39
                                                                                                                                                                                                                                                                                                                                                 ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: antisense sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 2
US-09-377-310-1/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-377-310-18
   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
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                                                                                                                     January 12, 2003, 09:46:10; Search time 47 Seconds (without alignments) 130.501 Million cell updates/sec
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Sequence 3,
Sequence 3,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sued_Patents NA:*
/cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/RD_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
                   GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-08-477-329-1

US-08-980-400-1

US-09-583-459A-1

US-09-583-459A-1

US-09-583-459A-1

US-09-683-459A-1

US-09-435-059-1

US-08-460-490-1

US-08-460-490-1

US-08-460-490-1

US-08-461-917-3

US-08-461-917-3

US-08-61-527-112

US-08-61-527-112

US-08-61-527-112

US-08-961-527-112

US-08-961-527-13
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US-09-377-310-38
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                       OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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seq length: 200000000
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20
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Match Length
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Result No.

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Minimum DB Maximum DB

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                                                                                                                                                                                                    APPLICATION NUMBER: US 07/146,877
APPLICATION NUMBER: US 07/146,877
FILING DATE: 22-JAN-1988
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/347,291
FILING DATE: 02-MAY-1989
ATTORNEY/ACENT INFORMATION:
NAME: Maki J.D., David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 990008.446
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ISSUE: 9
PAGES: 823-830
DATE: 1993-11
                                                                                                 TELEFAX: 206-682-6031
TELEX: 3723836
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JOURNAL: DN
VOLUME: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (233)..(3391)
PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DATABASE ACCESSION NUMBER: L13616/Genbank DATABASE ENTRY DATE: 1995-01-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDITION OF THE STREET: SCATTLE CITY: Seattle CITY: WA
                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
HYPOTHETICAL: N
                   MOLECULE TYPE:
                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                   TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 TTTCAACCAGATGGTCATTC 20
                                 POPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
                                                                                 ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20;
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                                                                 nucleic acid
                                                                                 4465 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6300 Columbia Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bell, Lillian A.

Kindsvogel Ph.D., Wayne R.

KINDSVOGEN PRODUCING SECRETED RECEPTOR ANALOGS
VENTION: MAID BIOLOGICALLY ACTIVE DIMERIZED POLYPEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sledziewski Ph.D., Andrzej Z
                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seed and Berry
                                                                                                                                                                        206-622-4900
                   CDNA
                                                   double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
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                                                                                                                                                                                                                                                                                                                                                                             US/07/634,510
                                                                                                                                                                                                                                                                                                                                                                                                                                                US/08/180,195
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                                                                                                                                                                                                          990008.446C3
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Pred. No. 0.28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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NAME/KEY: CDS;
LOCATION: 354.3671;
OTHER INFORMATION:
US-08-180-195-1
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Best Local Similarity 90.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/08477329 Patent No. 5750375
                                                                                                                                                                                                                   TELEFAX: 206-682-6031
TELEX: 3723836
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.24 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1263 TTTCATCCTGATGGTCATTC 1244
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Seed and
                                                                         ANTI-SENSE: N
ORIGINAL SOURCE:
                                                                                           HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORIGINAL SOURCE:
                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 4465 base pair
                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Bell, Lillian A. APPLICANT: Kindsvogel Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IMMEDIATE SOURCE:
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                                                                                                                       MOLECULE TYPE:
MMEDIATE SOURCE:
         ORGANISM: Homo sapiens
DEVELOPMENTAL STAGE: Adult
TISSUE TYPE: Skin
CELL TYPE: fibroblasts
                                                                                                                                                                                                                                    TELEPHONE: ZUU C
TELEPHONE: ZUU C
TERAX: 206-682-6031
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CELL TYPE: fibroblasts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DEVELOPMENTAL STAGE: Adult
TISSUE TYPE: Skin
                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 31, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLONE: pR-rX1
                                                                                                                                        TOPOLOGY:
                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM:
                                                                                                                                                                       1: 4465 base pairs nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Seed and Berry
6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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                                                                                                                                        linear
                                                                                                                                                                                                                                                                    206-622-4900
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07-JUN-1995
N. /^~
                                                                                                                            CDNA
                                                                                                                                                        double
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90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS AND BIOLOGICAL
                                                                                                                                                                                                                                                                                                             31,392
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                                                                                                                                                                                                                                                                                                   990008.44606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 16.8;
Pred. No. 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wayne R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
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RESULT 6
US-08-980-400-1/c
US-08-980-400-1/c

Sequence 1, Application US/08980400

Fatent No. 6018026

APPLICANT: Sledziewski Ph.D., Andrzej Z

APPLICANT: Bell, Lillian APPLICANT: Bell, Lillian APPLICANT: APPLICANT: Mindsvogel Ph.D., Wayne R.

TITLE OF INVENTION: METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS AND BIOLOGICAL

NUMBER OF SEQUENCES: 36

CORRESPONDENCE ADDRESS:

ADDRESSEE: Seed and Berry

STREET: 6300 Columbia Center, 701 Fifth Avenue

CITY: Seattle
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 Gaps
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 Indels
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90.0%; Pred. No. 13;
iive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIPCATION:
PRIOR APPLICATION:
PRIOR APPLICATION NUMBER: US 08/477,329
FILING DATE: 07-UW-1995
ATTORNEY/AGENT INFORMATION:
NAME: MAKE, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 990008.446C6
TELEPHONE: 206-622-4900
TELEPHONE: 206-682-6031
 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/980,400
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HYPOTHETICAL: N
ANTI-SENSE: N
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
DEVELOPMENTAL STAGE: Adult
TISSUE TYPE: Skin
                                                                        Db 1263 Tricarccreardercarrc 1244
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                                       1 TTTCAACCAGATGGTCATTC 20
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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 Conservative
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Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; NAME/KEY: CDS
; LOCATION: 354..3
; OTHER INFORMATION
US-08-980-400-1
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IMMEDIATE SOURCE:
CLONE: pR-rX1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
18;
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Matches
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APPLICANT: Bell, Lillian A.
APPLICANT: Bell, Lillian A.
APPLICANT: Kindsvogel Ph.D., Wayne R.
TITLE OF INVENTION: METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS AND BIOLOGICAL
NUMBER OF SROUBNCES: 36
CORRESPONDENCE ADDRESS:
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                                                                                                                                                  Gaps
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                                                                                                             DB 1; Length 4465;
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                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AUDRESSEE: Seed and Berry STREET: 6300 Columbia Center, 701 Fifth Avenue CITY: Seattle
                                                                                                         Query Match 84.0%; Score 16.8; D
Best Local Similarity 90.0%; Pred. No. 13;
Matches 18; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16.8; I
Pred. No. 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31,392
ama: 990008.446D5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTER: USA.

ZIP: 98104-7092

COMPUTER READABLE FORM:
MEDTUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHION DATA:
APPLICATION NUMBER: US/08/475,458
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MAKI, David J.
REGISTRATION NUMBER: 39,90008.446I
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-622-4900
TELEX: 3723836
INFORMATION POR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 4465 base pairs
"WUEF: MULTING ACID
                                                                                                                                                                                                                                                                           RESULT 5
US-08-475-458-1/c
Sequence 1, Application US/08475458
Patent No. 5843725
GENERAL INFORMATION:
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DEVELOPMENTAL STAGE: Adult
                                                                                                                                                                                     1 TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  84.0%;
90.0%;
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CELL TYPE: fibroblasts
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: CDNA
HYPOTHETICAL: N
ANTI-SENSE: N
             ; NAME/KEY: CDS
; LOCATION: 354..36
; OTHER INFORMATION:
US-08-477-329-1
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COTHER INFORMATION
US-08-475-458-1
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLONE: pR-rX1
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US-09-583-459A-1/c
                                                                                                                           US-09-583-459A-1
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                                                    Query Match 84.0
Best Local Similarity 90.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Sledziewski Ph.D., Andrzej
APPLICANT: Bell, Lillian A.
APPLICANT: Kindsvogel Ph.D., Wayne R.
TITLE OF INVENTION: METHODS OF PRODUCT
TITLE OF INVENTION: AND BIOLOGICALLY J
TITLE OF INVENTION: FUSIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 07/146,877
FILING DATE: 22-JAN-1988
APPLICATION UNMBER: US 07/347,291
FILING DATE: 02-MAY-1989
ATTORNEY/AGENT INFORMATION:
NAME: Maki J.D., David J.
REGISTRATION NUMBER: 31,392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 206-682-6031
TELEX: 3723836
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL: N
ANTI-SENSE: N
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                               TOPULUE TYPE: C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 99
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-622-4900
TELEFAX: 206-682-6031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/634,510
FILING DATE: 27-DEC-190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                     FEATURE:
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                                                                                                                                                                                                                                     MMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 4465 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                          NAME/KEY: CDS
LOCATION: 354..3671
OTHER INFORMATION:
                                                                                                                                                                                                                                                       TISSUE TYPE: Skin
CELL TYPE: fibroblasts
                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
DEVELOPMENTAL STAGE: Adult
TISSUE TYPE: Skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 30-MA CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
1 TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RY: USA
98104-7092
                                                                                                                                                                                                                  pR-rX1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: Seed and Berry
6300 Columbia Center
                                                                                                                                                                                                                                                                                                                                                                                     CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS AND BIOLOGICALLY ACTIVE DIMERIZED POLYPEPTIDE FUSIONS
                                                                     84.0%;
90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US/09/583,459A
                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     990008.446C3
                                                                     Score 16.8;
Pred. No. 13;
                                                        Mismatches
                                                                                          DB 4;
                                                                                      Length 4465;
                                                        0
                                                        Gaps
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US-09-583-210-1
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Query Match
Best Local Similarity 90.0
Matches 18; Conservative
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Patent No. 6291646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 07/634,510
FILING DATE: 27-DEC-1990
APPLICATION NUMBER: US 07/146,877
FILING DATE: 22-JAN-1988
APPLICATION NUMBER: US 07/347,291
FILING DATE: 02-MAY-1989
ATTORNEY/AGENT INFORMATION:
NAME: Maki J.D., David J.
REGISTRATION NUMBER: 31,392
REGISTRATION NUMBER: 990008.446
                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 206-682-6031
TELEX: 3723836
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Sledziewa...
APPLICANT: Bell, Lillian A.
APPLICANT: Bell, Lillian A.
APPLICANT: Kindsvogel Ph.D., Wayne R.
APPLICANT: KINDSVOGEN AND METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS
TITLE OF INVENTION: AND BIOLOGICALLY ACTIVE DIMERIZED POLYPEPTIDE
TITLE OF INVENTION: FUSIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1263 TTTCATCCTGATGGTCATTC 1244
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/583,21
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                    FEATURE:
                                                                                                                                                                                        IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                           ANTI-SENSE: N
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 4465 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 206-622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
                                                                                          LOCATION: 354..36
OTHER INFORMATION:
                                                                                                             NAME/KEY: CDS
LOCATION: 354..3671
                                                                                                                                                                                                       TISSUE TYPE: Skin
CELL TYPE: fibroblasts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 3
CLASSIFICATION:
                                                                                                                                                                     CLONE:
                                                                                                                                                                                                                                         ORGANISM: Homo sapiens
DEVELOPMENTAL STAGE: A
                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                      double
 84.0%; Score 16.8; 1
90.0%; Pred. No. 13;
tive 0; Mismatches
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                                                                                                                                                                                                                                             Adult
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    990008.446C3
                                       DB 4;
     2
                                       Length 4465;
     <u>.</u>
     Gaps
     0,
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TTTCAACCAGATGGTCATTC 20

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RESULT 10
US-09-435-059-1/C
Sequence 1, Application US/09435059
; Sequence 1, Application US/09435059
; Patent No. 6323323
; GENERAL INFORMATION:
; APPLICANT: Sledatewski Ph.D., Andrzej Z
APPLICANT: Rindsvogel Ph.D., Wayne R.;
; TITLE OF INVENTION: METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS AND BIOLOGICAL
; VUMBER OF SEQUENCES: 36
; CORRESPONDENCE ADDRESS:
; ADDRESSES: Seed and Berry
; STREET: 6300 Columbia Center, 701 Fifth Avenue
; CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/435,059
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477,329
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: MAXI, DAVIG J.
REGISTRATION NUMBER: 31,392
REFRENCY/DOCKET NUMBER: 990008.446C6
TELECOMMULICATION INFORMATION:
TELEPHONE: 206-682-4900
TELEPHONE: 206-682-4900
TELEPHONE: 246-6031
TELEPHONE: 246-6031
TELEPHONE: 4465 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TYPE: nucleic acid
STRANDEDNESS: double
TOPOGLOGY: linear
TYPE: nucleic acid
STRANDEDNESS: double
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90.0%; Pred. No. 13;
Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: WA
COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPACIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                               1263 TTTCATCCTGATGGTCATTC 1244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
DEVELOPMENTAL STAGE: Adult
      1 TITCAACCAGATGGTCATIC 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; LOCATION: 354..36
; OTHER INFORMATION:
US-09-435-059-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 18; Conserv
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CELL TYPE: fibr
IMMEDIATE SOURCE:
CLONE: pR-rX1
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US-09-581-449A-1/c

; Sequence 1, Application US/09583449A
; Patent No. 6300099
; GENERAL INFORMATION:
; APPLICANT: Sledziewski Ph.D., Andrzej Z
; APPLICANT: Bell, Lillian A.
; APPLICANT: Rell, Lillian A.
; TITLE OF INVENTION: METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS
; TITLE OF INVENTION: AND BIOLOGICALLY ACTIVE DIMERIZED POLYPEPTIDE
; TITLE OF INVENTION: FUSIONS
; NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSE: Seed and Berry
STREET: 6300 Columbia Center
CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 98104-7092
ZIP: 98104-7092
ZIP: 98104-7092
ZIP: 98104-7092
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/583,449A
FILING DATE: 27-DEC-1990
APPLICATION NUMBER: US 07/46,877
FILING DATE: 27-DEC-1990
APPLICATION NUMBER: US 07/34,510
FILING DATE: 02-MAY-1989
ATTORNEY/AGENT INFORMATION:
NAME: MAXI JO.D., David J.
RECISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 390008.446C
TELLEPAX: 206-682-6031
FILING DATE: 3723836
TELLEPAX: 3723836
INFORMATION FOR SEQ ID NO: 1:
ENGTH: 4465 base pairs
INFORMATION FOR SEQ ID NO: 1:
ENGTH: 4465 base pairs
INFORMATION FOR SEQ ID NO: 1:
FIRM STRANDEDNES: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99000B.446C3
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DEVELOPMENTAL STAGE: Adult
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CELL TYPE: Skin
CELL TYPE: fibroblasts
IMMEDIATE SOURCE:
CLONE: PRATURP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: CDS
LOCATION: 354..3671
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: cDNA HYPOTHETICAL: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ANTI-SENSE: N
ORIGINAL SOURCE:
ORGANISM: HOMC
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US-08-168-917-1/c
                                                                    US-08-460-510-1/c
                                                                                    RESULT 12
                                                                                                                                                                                                                                                                           NAME/KEY:
LOCATION:
US-08-168-917-1
              Sequence 1, Application US/08460510 Patent No. 5872218 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Appli
Patent No. 568657
                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNE.... Ching, Edwl: 5.4,090
REGISTRATION NUMBER: 34,090
REFERENCE/DOCKET NUMBER: 1241f
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 11
SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pairs
TYPE: nucleic acid
                                                                                                                                        1096 TTTCATCCTGATGGTCATTC 1077
APPLICANT: Wolf, David
                                                                                                                                                                                                                                                                                                                                                                             ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Escobedo, Jaime A. APPLICANT: Williams, Lewis T.
                                                                                                                                                                                                                                                                                                                                                                                                   HYPOTHETICAL: 1
                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGICAL: CD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: DOMAINS OF EXTRACELLULAR REGION OF HUMAN
TITLE OF INVENTION: PLATELET-DERIVED GROWTH FACTOR RECEPTOR POLYPEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo Sapiens
STRAIN: lambda gt10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                        1 TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94105
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fretto, Larry J
Giese, Neill A.
                                                                                                                                                                                                            Conservative
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Tomlinson, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                   cDNA to mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     double
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Larry J.
                                                                                                                                                                                                                        84.0%;
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                                                                                                                                                                                                                          Score 16.8;
Pred. No. 13;
                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                          Length 5427;
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                                                                                                                                                                                                              Gaps
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RESULT 13
US-08-460-490-1/c
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                                                                                                                                                                                                                  Sequence 1, Application US/08460490 Patent No. 5891652
                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                          GENERAL INFORMATION:

APPLICANT: Wolf, David

APPLICANT: Tomlinson, James E.

APPLICANT: Fretto, Larry J.

APPLICANT: Giese, Neill A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (415) 326-24: INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                             1096 TTTCATCCTGATGGTCATTC 1077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.
CURRENT APPLICATION DATA:
TT/(NO)/46 E10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: Une ....
STREET: San Francisco
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/460,510 FILING DATE: 02-JUN-1995 CLASSIFICATION: 435 ATTORNEY/AGENT INFORMATION:
                                                    APPLICANT: Williams TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Williams
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Dow, Karen B.
REGISTRATION NUMBER: 29,684
REFERENCE/DOCKET NUMBER: 01
                                                                                                                                                                                                                                                                                                                                                                                         84.0%;
Local Similarity 90.0%;
es 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN: lambda gt10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                             1 TTTCAACCAGATGGTCATTC 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94105
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One Market Plaza, Steuart Street Tower, Suite 2000
                                                                                    Escobedo, Jaime A. Williams, Lewis T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Escobedo, Jaime A.
Williams, Lewis T.
WILLIAMS, LOWAINS OF EXTRACELLULAR REGION OF HUMAN
VENTION: PLATELET-DERIVED GROWTH FACTOR RECEPTOR POLYPEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (415) 326-2422
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Giese, Neill A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tomlinson, James E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             187..3504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
TOWNSEND and TOWNSEND and CREW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sapiens
                                                      DOMAINS OF EXTRACELLULAR REGION OF HUMAN PLATELET-DERIVED GROWTH FACTOR RECEPTOR POLYPEPTIDES
                                     23
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 5427;
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0; Mismatches
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90.0%; Pred. No. 13
                                              FILING DATE: 10-FEB-1989

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/151,414
FILING DATE: 02-FEB-1988
ATTORNEY/AGENT INFORMATION:
NAME: Dow, Karen B.
REGISTRATION NUMBER: 29,684
REFERENCE/DOCKET NUMBER: 2307K-267-2
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: 415/326-242
INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pairs
TYPE: nucleic acid
STRANDENNESS: single
TOPPLOGY: linear
MOLECH F TOWE: ON NO: 10 NO: 
                         APPLICATION NUMBER: US 07/309,322
FILING DATE: 10-FEB-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 3, Application US/08461917
; Patent No. 6372438
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REGISTRATION NUMBER: 29,684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 23
TELECOMMUNICATION:
TELEPHONE: 415/326-2400
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Best Local Similarity 90.0'
Matches 18; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
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CITY: San Francisco
STATE: California
COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY:
LOCATION:
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US-08-461-917-3/C
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San Francisco
California
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GENERAL INFORMATION:

APPLICANT: WILLIAMS, Lewis T.

APPLICANT: ESCOBEDO, Jaime A.

TITLE OF INVENTION: PLATELET-DERIVED GROWTH FACTOR RECEPTORS

NUMBER OF SEQUENCES: 14

CORRESPONDENCE ADDRESS:

ADDRESSEE: Townsend and Townsend and Crew
STREET: One Market, Steuart Street Tower, 20th Floor

CITY: San Francisco

STATE: California

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       84.0%; Score 16.8; DB 2; Length 5427; 90.0%; Pred. No. 13;
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                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,490
FILING DATE: 02-UN-1995
CLASSITCATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Dow, Karen B.
REGISTRATION NUMBER: 012418-001420
TELEFROCHOCKET NUMBER: 012418-001420
TELEFROCHOCKET NUMBER: 012418-001420
TELEFROCHOIN INFORMATION:
TELEFROCHONE: (415) 326-240
TELEFROCH (415) 326-242
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 5427 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
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FILING DATE: 05-UTN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 94.05
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3, Application US/08462728
Patent No. 6043211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1096 rrrcarccrgargercarrc 1077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: CDNA to mRNA HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo Sapiens
STRAIN: lambda gt10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 90.0
Matches 18; Conservative
                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ) NAME/KEY: CDS
; LOCATION: 187..3504
US-08-460-490-1
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ORIGINAL SOURCE:
                                                                                                                                                   94105
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Gaps
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APPLICANT: WILLIAMS, Lewis T.
APPLICANT: ESCOBEDO, Jaime A.
TITLE OF INVENTION PLATELET-DERIVED GROWTH FACTOR RECEPTORS
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
DB 3; Length 5427;
                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Townsend and Townsenu and Treet Tower, 20th Floor STREET: One Market, Steuart Street Tower, 20th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CUMINER: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,917
FILING DATE: 05-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 07/309,322
FILING DATE: 10-FEB-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/151,414
FILING DATE: 02-FEB-1988
ATFORDEY AGENT INFORMATION:
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Sequence 9537, Apples Sequence 3, Appli Sequence 242, Apples Sequence 248, Apples Sequence 180, Apples Sequence 111, Apples Sequence 28, Apples Sequence 975, Apple

Sequence 712, App Sequence 2112, App Sequence 1609, Ap Sequence 3, Appli Sequence 21937, A Sequence 3281, Ap Sequence 14297, Ap

Sequence

Sequence:

Run on:

Searched:

Database

Result Š.

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US-09-757-100B-18

Sequence 18, Application US/09757100B

Sequence 18, Application US/09757100B

Sequence 18, Application US/09757100B

Sequence 18, Application Garde, William A.

APPLICANT: Garde, William A.

APPLICANT: Nero, Pamela S.

TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase

TITLE OF INVENTION: Expression

FILE REFERENCE: ISPH-053

CURRENT FILING DATE: 2001-03-15

PRIOR PILING DATE: 1999-08-19

PRIOR FILING DATE: 1999-08-19

PRIOR FILING DATE: 2000-07-13

NUMBER OF SEQ ID NOS: 44

SOFTWARE: PETENTIN Ver. 2.0

SEQ ID NO. 18
0. US-09-815-242-9137

0. US-09-915-242-9537

0. US-09-938-842A-3191

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0. US-09-938-842A-3191

0. US-09-974-300-180

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0. US-09-9138-626-1

0. US-09-10-188-171

0. US-09-175-100B-38

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0. US-09-9864-761-21937

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0. US-09-864-761-21937

0. US-09-864-761-2203

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0. US-09-864-761-2203

0. US-09-864-761-2203
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100.0%; Pred. No. 0.13;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                  ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: antisense sequence US-09-757-100B-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 981, Application US/09736457
Patent No. US20020168637A1
PAPELI INFORMATION:
APPLICANT: Wang, Tongtong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20
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ORGANISM: Artificial Sequence
    TTTCAACCAGATGGTCATTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TTTCAACCAGATGGTCATTC
                                                                                                                         3309400
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4187
41104
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Best Local Similarity
Matches 20; Conserv
  RESULT 2
US-09-736-457-981
    셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 981, Apple Sequence 11, Appli Sequence 21874, Ap Sequence 5154, Ap Sequence 23, Appl Sequence 13, Appl Sequence 15, Appl Sequence 15, Appl Sequence 17, Appl Sequence 17, Appl Sequence 19, Appl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15, Appl
Sequence 17, Appl
Sequence 19, Appl
Sequence 21, Appl
Sequence 1, Appl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18, Appl
Sequence 981, App
Sequence 981, App
                                                                                       January 12, 2003, 10:38:40; Search time 56 Seconds (without alignments) 157.180 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Sequence 1678
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1: /cgn2_6/ptodata/2/pubpna/UGO7_PUBCOMB.seq:*
2: /cgn2_6/ptodata/2/pubpna/PCT7_NW_PUBL.seq:*
2: /cgn2_6/ptodata/2/pubpna/UGO6_PUBCOMB.seq:*
4: /cgn2_6/ptodata/2/pubpna/UGO6_PUBCOMB.seq:*
5: /cgn2_6/ptodata/2/pubpna/UGO6_PUBCOMB.seq:*
6: /cgn2_6/ptodata/2/pubpna/UGO6_PUBCOMB.seq:*
7: /cgn2_6/ptodata/2/pubpna/UGO8_NEW_PUB.seq:*
8: /cgn2_6/ptodata/2/pubpna/UGO8_NEW_PUB.seq:*
9: /cgn2_6/ptodata/2/pubpna/UGO9_PUBCOMB.seq:*
10: /cgn2_6/ptodata/2/pubpna/UGO9_PUBCOMB.seq:*
11: /cgn2_6/ptodata/2/pubpna/UGO9_PUBCOMB.seq:*
13: /cgn2_6/ptodata/2/pubpna/UGO9_PUBCOMB.seq:*
13: /cgn2_6/ptodata/2/pubpna/UGO0_PUBCOMB.seq:*
14: /cgn2_6/ptodata/2/pubpna/UGO0_PUBCOMB.seq:*
14: /cgn2_6/ptodata/2/pubpna/UGO0_PUBCOMB.seq:*
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               GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                                                                                           389086 seqs, 220051671 residues
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                                                                 OM nucleic - nucleic search, using sw model
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 2000000000
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Match Length DB
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Length 20;

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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER FILE REFERENCE: 210121.478C15
CURRENT APPLICATION NUMBER: US/09/736,457
CURRENT FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 1864
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 981
LENGTH: 550
TYPE: DNA
ORGANISM: Homo sapien
US-09-736-457-981
US-09-849-626-981; Sequence 981, App.; Publication No. 1
                                                                                                                                                                                                                                                         CURRENT APPLIATION NUMBER: US/09/902,941
CURRENT FILING DATE: 2001-07-10
NUMBER OF SEQ ID NOS: 2002
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 981
LENGTH: 550
TYPE: DNA
ORGANISM: Homo sapiens
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APPLICANT: Bengur, Chaitanya S.
APPLICANT: McNabb, Andria
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.478C17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Henderson, Robert A.
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                                                                                                                                               1 TTTCAACCAGATGGTCATTC 20
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                                                                                                               TTTCAACCAGATGGTCATTC 214
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Johnson, Jeffrey C.
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Watanabe, Yoshihiro
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             Application US/09849626
No. US20020197669A1
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Carter, Darrick
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Fan, Liqun
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                                                                                                                                                                              100.0%; Score 20; DB 9; Length 550; ilarity 100.0%; Pred. No. 0.26; Conservative 0; Mismatches 0; Indels
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Pred. No. 0.26;
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FILE REFERENCE: ISPH-0533
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 1999-08-19
PRIOR APPLICATION NUMBER: PCT/US00/18999
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEO ID NOS: 44
SOFTWARE: PALENTIN Ver. 2.0
SEQ ID NO 1
LENGTH: 3791
                                                                                                                                                 US-09-757-100B-1
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                                                                                                                                                             NAME/KEY: CDS
LOCATION: (233)..(3391)
PUBLICATION INFORMATION:
JOURNAL: DNA
VOLUME: 12
ISSUE: 9
PAGES: 823-830
PAGES: 823-830
PATS: 1993-11
DATABASE ACCESSION NUMBER: L13616/Genbank
DATABASE ENTRY DATE: 1995-01-02
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CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 1926
SOFTWARE: FREESEQ for Windows Version 3.0
SEQ ID NO 981
LENGTH: 550
                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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Patent No. US20010034329A1
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APPLICANT: Gaarde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Switzer, Anne
APPLICANT: McNeill, Patricia
APPLICANT: Clapper, Jonathan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.478C16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Bangur, Chaitanya
APPLICANT: Fanger, Gary
APPLICANT: Wang, Aijun
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                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                         FEATURE:
3529
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Local Similarity 100.0%;
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                           TTTCAACCAGATGGTCATTC 20
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                                                                     Conservative
                                                                                       100.0%; Score 20; DB 10; 100.0%; Pred. No. 0.39;
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Pred. No. 0.26;
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US-864-761-1514

Sequence 5154, Application US/09864761

Sequence 5154, Application US/09864761

Sequence 5154, Application US/09864761

Sequence 5154, Application US/09864761

Sequence 5154, Application General Control Co
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OTHER INFORMATION: MAP TO AL121654.1
OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 0.97
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 0.84
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Annomax Sequence Listing Engine vers. 1.1
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SOFTWARE: Annomax Sequence I
SEQ ID NO 5154
LENGTH: 400
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Best Local Similarity
Matches 18; Conserv
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OTHER INFORMATION: EXPRESSED IN BY474, SIGNAL = 0.97

OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 0.84

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1

OTHER INFORMATION: EXPRESSED IN BONE WARROW, SIGNAL = 1

OTHER INFORMATION: SWISSPROT HIT: P47144, EVALUE 3.50e-02
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SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 21875
LENGTH: 244
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90.0%; Pred. No. 10;
iive 0; Mismatches 2;
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Best Local Similarity 90.0
Matches 18; Conservative
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ORGANISM: Homo sapiens
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OTHER
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; OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL =
; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL
US-09-864-761-16204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-864-761-16204
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16204, Applicatio Patent No. US20020048763A1
                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 09/632,366 PRIOR FILING DATE: 2000-08-03
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                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                              ENGTH: 528
344
                                                                                                   Local Similarity
                                      1 TTTCAACCAGATGGTCATTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 60/236,359 FILING DATE: 2000-09-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: PCT/US01/00670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 09/774,203
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 09/608,408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: PCT/US01/00662
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FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 60/234,687
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TTTCAACCAGATGTACATTC 363
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Chen, Wensheng
                                                                                  Conservative
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                                                                                                   Score 16.8;
Pred. No. 12
                                                                             Mismatches
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                                                                                                                        Length 528;
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                                                                                  Gaps
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RESULT 9

US-09-866-510-15/c ; Sequence 15, Application US/09866510 ; Patent No. US20020111304A1

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; TYPE: DNA
; ORGANISM: Homo sapiens
; PEATURE:
; PEATURE:
; NAME/KEY: CDS
; LOCATION: (1)...(1683)
US-09-866-510-23
RESULT 11
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                                                                                                                        Best Loc
Matches
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Best Local Similarity
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                                                                                                                                                            Query Match
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/250,747
PRIOR FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: 60/289,103
PRIOR FILING DATE: 2001-05-07
PRIOR FILING DATE: 2001-05-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: KAZLAUSKAS, ANDRIUS
APPLICANT: IKUNO, YASUSHI
TITLE OF INVENTION: METHODS AND CÔMPOSITIONS FOR TREATING FIBROTIC DISEASES
FILE REFERENCE: ERM-104.01
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NUMBER OF SEQ ID NOS: 33
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CURRENT APPLICATION NUMBER: US/09/866,510
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/250,747
PRIOR FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: 60/289,103
                                                                                                                                                                                                                                                                                                                                            SOFTWARE:
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TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING FIBROTIC DISEASES
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APPLICANT: IKUNO, YASUSHI
                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 33
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                                                                                                                                                                                                                                                   ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                               NAME/KEY: CDS
LOCATION: (1)..(3270)
                                                                                                                                                                                                                                                                                         TYPE: DNA
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18; Conserv
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                                                                                                                          Conservative
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Pred. No. 18;
0; Mismatches
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Pred. No. 1
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Sequence 21, Application US/09866510

Patent No. US20020111304A1

GENERAL INFORMATION:

APPLICANT: KAZLAUSKAS, ANDRIUS

APPLICANT: KAZLAUSKAS, ANDRIUS

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING FIBROTIC DISEASES

FILE REFERENCE: ERM-104.01

CURRENT APPLICATION NUMBER: US/09/866,510

CURRENT FILING DATE: 2001-05-25

PRIOR FILING DATE: 2000-12-01

PRIOR FILING DATE: 2000-12-01

PRIOR FILING DATE: 2001-05-07

NUMBER OF SEQ ID NOS: 33

SOFTWARE: PATENTIN VOR: 33

SOFTWARE: PATENTIN VOR: 2.1

FENOM: 201
             INVENTION: METHODS AND COMPOSITIONS FOR TREATING FIBROTIC DISEASES
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Sequence 1, Application US/09955363
Patent No. US20020173621A1
GENERAL INFORMATION:
BELLICANT: Sledziewski Ph.D., Andrzej Z
APPLICANT: Bell, Lillian A.
Kindsvogel Ph.D., Wayne R.
TITLE OF INVENTION: METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS
AND BIOLOGICALLY ACTIVE DIMERIZED POLYPEPTIDE
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CURRENT APPLICATION NUMBER: US/09/866,510
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/250,747
PRIOR FILING DATE: 2001-05-05
PRIOR FILING DATE: 2000-12-01
PRIOR FILING DATE: 2000-12-03
NUMBER OF SEQ ID NOS: 33
SOFTWARE: PALENTIN Ver. 2.1
SEQ ID NO 19
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90.0%; Pred. No. 18;
iive 0; Mismatches
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Best Local Similarity 90.0%; Pred. No. 18;
Matches 18; Conservative 0; Mismatches
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Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                  ) NAME/KEY: CDS
; LOCATION: (1)..(3318)
US-09-866-510-19
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US-09-866-510-21/c
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LOCATION: (1)
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US-09-955-363-1/c
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                                              APPLICANT: IKUNO, YASUSHI
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING FIBROTIC DISEASES
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING FIBROTIC DISEASES
CURRENT APPLICATION NUMBER: US/09/866,510
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/280,747
PRIOR APPLICATION NUMBER: 60/289,103
PRIOR PILING DATE: 2000-12-01
PRIOR FILING DATE: 2000-05-07
NUMBER OF SEC ID NOS: 33
SOFTWARE: Patentin Ver. 2.1
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APPLICANT: KAZLAUSKAS, ANDRIUS
APPLICANT: KAZLAUSKAS, ANDRIUS
TITLE OF INVENTION METHODS AND COMPOSITIONS FOR TREATING FIBROTIC DISEASES
FILE REFERENCE: ERM-104.01
CURRENT APPLICATION NUMBER: 06/255
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 66/250,747
PRIOR PILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: 66/289,103
PRIOR PILING DATE: 2001-05-07
NUMBER OF SEQ ID NOS: 31
SOFTWARE: PATENTIN VET. 2.1
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Best Local Similarity 90.0%; Pred. No. 18
Matches 18; Conservative 0; Mismatche
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US-09-866-510-17/c
; Sequence 17, Application US/09866510
; Patent No. US20020111304A1
; GENERAL INFORMATION:
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GENERAL INFORMATION:
APPLICANT: KAZLAUSKAS, ANDRIUS
APPLICANT: IKUNO, YASUSHI
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                            ; NAME/KEY: CDS
; LOCATION: (1)..(3318)
US-09-866-510-15
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; LOCATION: (1)..(3318)
US-09-866-510-17
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                                                                                                                                                                                                                                                                                              SEQ ID NO 15
LENGTH: 3321
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STATE: WA
COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARR: PACENTIN Release #1.24
CURRENT APPLICATION NUMBER: US/09/955,363
FILING DATE: 18-Sep-2001
CLASSIFICATION NUMBER: US 07/634,510
PRIOR APPLICATION NUMBER: US 07/634,510
PRIOR APPLICATION NUMBER: US 07/347,291
FILING DATE: <Unknown>
APPLICATION NUMBER: US 07/347,291
FILING DATE: 02-MAY-1989
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 990008.446C3
TELEPHONE: 206-622-4900
TELEPHONE: 206-622-4900
TELEFAX: 372336
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 4465 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: N
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
DEVELOPMENTAL STAGE: Adult
TISSUE TYPE: fibroblasts
IMMEDIATE SOURCE:
CLONE: pR-rX1
                                                                                                                                                                                                                          ; INAME/KEY: CDS; LOCATION: 354..3671; OTHER INFORMATION: SEQUENCE DESCRIPTION: SEQ ID NO: 1: US-09-955-363-1
Search completed: January 12, 2003, 12:08:34 Job time : 59 secs
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                                                                                                                                                      Query Match 84.0%; Score 16.8; I Best Local Similarity 90.0%; Pred. No. 19; Matches 18; Conservative 0; Mismatches
                                                                            1263 TTTCATCCTGATGGTCATTC 1244
                                                                                                                     1 TTTCAACCAGATGGTCATTC 20
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CORRESPONDENCE ADDRESSE: Seed and Berry
STREET: 6300 Columbia Center
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STATE: WA
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Sequence:

Minimum DB Maximum DB

Database

Searched:

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BF669338 602120623
AW992843 RC2-BN003
AW992843 RC2-BN003
AW999847 RC2-BN003
AQ42426 CTTB1-E1-
BF97799 60214726
BF5756 601340992
BF701452 602125760
BF701452 602125760
BF701452 602125760
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BF77595 602119853
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BF217100 601892483
BF2171100 601892483
BF7171101 60181331
              AA883514 Or41g10.s
BM664187 UI-B-CII-
NA230 yw21h06.r1
A1341506 gx92a12.x
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BG986106 PM2-DT004
A1016118 oc134g97.s
AA404694 zt41a11.r
AA80178 tf98e02.x
BM790849 K-EST0070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 28s)

Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deoliveire, F.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
                                                                                                                                                                                                                                                                                                         BF213328 601845047
BG110386 602279286
BG617438 602615284
B1917958 603181742
                                                                                                                                                                                                                                                                                                                                                                                                         BF997449 28- 285 bp mRNA linear EST 23-JAN-2001
CMO-GN0100-081100-576-a05 GN0100 Homo sapiens cDNA, mRNA sequence.
                                                                                                     AA453479 zx47a09.s
AW003490 wq66e03.x
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20202663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rud Prof. Antonio Prudente 109, 4 andar, 01509-010,
Brazil
                                                                                                                                                                                                                                                                                                                                                                ALIGNMENTS
                                                                                                              AW003490
BF669338
AW992843
AG060293
BF691197
                                                                                                                                                         AW992847
AQ424326
BF701453
BE567556
BQ438629
AQ438629
BQ565639
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BG517048
                AA983514
BM664187
N42230
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BG986106
AI016118
AA404694
AI380178
BM790849
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BF997449.1 GI:12403772
0110
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                                                                                                               Tel: +55-11-2704922
Fax: +55-11-2707001
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RESULT 1
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AUTHORS
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SOURCE
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R32551 yh54d07.r1
AV709841 AV709841
BE092864 CM2-BT074
R89170 yq02b06.r1
AV724065 AV724065
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                                                          ; Search time 1993 Seconds (without alignments) 162.524 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Description
                                                                                                                                                                           32308132
       GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                         Total number of hits satisfying chosen parameters:
                                                                                                                                                         16154066 seqs, 8097743376 residues
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                                          - nucleic search, using sw model
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AV709841
BE092864
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Match 100%
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em_gss_pro:*
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seq length: 200000000
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gb_htc:*
gb_est3:*
gb_est3:*
gb_est5:*
em_estfun:*
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Sao Paulo-SP,

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Result Š.

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JOURNAL
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AUTHORS
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R32551/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20;
                                                                                                                                                      Email: est@watson.wustl.edu
Insert Size: 691
High quality sequence stops: 299
Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Insert Length: 691
Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                 ,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.
The WashU-Merck EST Project
Unpublished (1995)
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=CMO&t2=CMO-GN0100-
081100-576-a05&t3=2000-11-08&t4=1)
Seq_primer: puc_l8 forward
                                                                                                                                        Seq
                                                                                                                                                                                                                                                                                                                                                                      Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R32551 351 bp mRNA linear EST 28-APR-19:
yh54d07.r1 Soares placenta Nb2HP Homo sapiens cDNA clone
YMAGE:133549 5' similar to gb:L05186 FOCAL ADHESION KINASE (HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence start: 10 High quality sequence stop: 285.
                                                                                                                                                                                                                                                                                                                           Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hillier, L., Clark, N., Dubuque, T., Elliston, K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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1 (bases 1 to 351)
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                                                                                                                                        primer: M13RP1
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                                                                                        quality sequence stop: 299
Location/Qualifiers
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/db_xref="GDB:539275"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="GN0100"
/dev_stage="Adult"
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                                             organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Homo sapiens"
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Pred. No. 21;
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173
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                                                         20;
                                                                                                                                                                                                                                                                                                                                                                        Contact: Zeguang Han
Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai
201203, P. R. China
                                                                                                                                                                                                                                                                                                                                                                                                                                  Peng,Y., Song,H., Huang,Q., Huang,C., Gu,Y., Yang,Y., Gao,H., Xu,X., Li,N., Qian,B., Liu,F., Qu,J., Gao,X., Cheng, Zeng,L., Xu,S., Gu,W., Tu,Y., Jia,J., Fu,G., Ren,S., Zhon,G., Hu,R., Chen,J., Chen,Z. and Han,Z. Homo sapiens cDNA ADC clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 392)
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AV709841.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AV709841 ADC Homo sapiens cDNA clone ADCAGA11 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                  Email: hanzg@chgc.sh.cn
This clone is available at CHGC
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EST
                                                                                                                                                                                                                                                                                                                                               Tel: 86-21-50801919(ex.45)
Fax: 86-21-50801922
                                                        Similarity 100.
20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not RI stress of the modified pTTT3 vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo. "

66 c 65 g 120 t
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/dev_stage="Adult"
/lab_host="SOLR"
                                                                                                                                                                                                                          /db_xref="taxon:9606"
/clone="ADCAGA11"
/clone_lib="ADC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="Soares placenta Nb2HP"
/sex="Female"
                                                                                                                                                                                                                                                                       organism="Homo sapiens"
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                                                                       100.0%;
                                                                                                                                                                e="Vector: pBluescript sk(-); Site_1: EcoRI; Site_2:
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Pred. No. 24;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammaliai Eutheria; Primates; Catarrhini; Hominidae; Homo.

E L (bases 1 to 416)

S Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston, R., Williamson,A., Wohldmann,P. and Wilson,R.

The Washlorder K EST Project

AL Unpublished (1995)

Contact: Wilson RK
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800

Fax: 314 286 1800

Email: est@watson.wustl.edu

Insert Size: 1206

High quality sequence stops: 337

Source: IMAGE Consortium, LLNL

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.

Insert Length: 1206 Std Error: 0.00

Seq primer: M13RPl

High quality sequence stops: 337.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 424)
Gu,Y., Peng,Y., Song,H., Huang,Q., Yang,Y., Gao,G., Xiao,H., Xu,X., Li,N., Qian,B., Liu,F., Qu,J., Gao,X., Cheng,Z., Xu,Z., Zeng,L., Xu,S., Gu,W., Tu,Y., Jia,J., Fu,G., Ren,S., Zhong,M., Lu,G., Hu,R., Chen,J., Chen,Z., and Han,Z.
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AV724065 AVE Homo sapiens cDNA clone HTBAIF11 5', mRNA sequence. AV724065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1. .416
/organism="Homo sapiens"
/db_xref="GDB:3764781"
/db_xref="taxon:9606"
/clone="INMGE:195731"
/clone_lib="Soares fetal liver spleen INFLS"
/sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 20; DB 14; 100.0%; Pred. No. 26;
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Best Local Similarity
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AV724065/c
LOCUS
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AUTHORS
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/db xref="taxon:9606"
/dbo=lib="BT0741"
/dev stage="Adult"
/dev stage="Adult"
/note="Organ: breast; Vector: puc18; Site_1: SmaI; Site_2: SmaI; A mini-library was made by cloning products derived from ORESTES FOR (U.S. Letters Patent application No. 196; 716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low
                                                                                                                                                                                                                        Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases I to 412)

1 bass I to 412)

1 bass Neto, E. Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Slmpson, D.H., Brunstein, A., deoliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and Simpson, A.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: +55-11-2704922
Fax: +55-11-2707001
Email: asimpson@ludwig.org.br
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=&t2=CM2-BT0741-050
400-140-c128t3=2000-04-05&t4=1)
Seq primer: puc 18 forward
High quality sequence start: 39
High quality sequence stop: 410.
                                                                       BE092864 11-050400-140-c12 BT0741 Homo sapiens cDNA, mRNA sequence.
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Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                Shotgun sequencing of the human transcriptome with ORF expressed
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Proc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)
20202663
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Pred. No. 26;
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Best Local Similarity 100.0%;
Matches 20; Conservative 0
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R89170
R89170.1 GI:953997
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Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai
201203, P. R. China
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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601339461F1 NIH_MGC_53 Homo
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Fax: 86-21-50801922
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High quality sequence stop: 421.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: hanzg@chgc.sh.cn
This clone is available at CHGC in Shanghai.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CDNA Library Preparation: CLONTECH Laboratories, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (bases 1 to 425)
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/clone_lib="NIH_MGC_53"
//clone_lib="NIH_MGC_53"
//tissue_type="carcinoma, cell line"
//tissue_type="carcinoma, cell line"
//lab host="UBIOB (TI phage-resistant)"
//note="Organ: bladder; Vector: pDNR-LIB (Clontech);
//note="Organ: bladder; Vect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ø
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/tissue_type="Hypothalamus"
/dev_stage="Adult"
/lab_host="SOLR"
                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="HTBAIF11"
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                                                                                                                                                                                                                                                                                                                                                                                       clone="IMAGE:3681878"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      90 c
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s cDNA clone
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IMAGE:3681878 5',
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AUTHORS
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AW136503
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TTTCAACCAGATGGTCATTC 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The sequence contained an oligo-dT track that was present in the oligonuclectide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide poly A tail cDNA Library Preparation: M.B. Soares Lab Clone distribution: NCI-CGAP clone distribution information can be found through the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Mammalia; Eutheria; 1 (bases 1 to 429)
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AW136503.1 GI:6140563
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seq primer: M13 Forward
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118 c 91 g 99 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone="IMAGE:2715763"
/clone_lib="NCI_CGAP_Sub3"
/lab_host="DH10B_(Life_Technologies)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                db_xref="taxon:9606"
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Pred. No.
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26;
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Gaps

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BASE COUNT ORIGIN

DEFINITION

RESULT 9 AA983514

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ACCESSION

VERSION KEYWORDS

ORGANISM

SOURCE

AUTHORS TITLE

JOURNAL

COMMENT

FEATURES

REFERENCE

BASE COUNT ORIGIN

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/closue_lib="lib"
/tissue_type="RPE and Choroid"
/dev stage="RPE and Choroid"
/dev stage="adult"
/lab_host="DH10B (Life Technologies) (Tl phage resistant)"
/note="Organ: eye; Vector: pTT3-Pac (Pharmacia) with a
modified polylinker; Site 1: ECOR 1; Site 2: Not 1;
UI-E-CI1 is a normalized CDNA library containing the
following tissue(s): RPE and Choroid. The library was
constructed according to Bonaldo, Lennon and Soares,
Genome Research, 6:791-806, 1996. First strand cDNA
synthesis was primed with an oligo-dT primer containing a
Not 1 site. Double stranded cDNA was ligated to an EcoR I
adaptor, digested with Not I, and cloned directionally
into pT/T3-Pac vector. The oligonucleotide used to prime
the synthesis of first-strand cDNA contains a library tag
sequence that is located between the Not I site and the
(dT):B tail. The sequence tag for this library is ACCTA.
This library was created for the program, Gene Discovery
in the Visual System, supported by National Eye Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       University of lowa University of lowa University of lowa University of lowa 451 Eckstein Medical Research Building lowa City, IA 52242, USA Tel: 319 315 9250
Fax: 319 315 9250
Fax: 319 315 9565
Email: msoares@blue.weeg.uiowa.edu
Tissue Procurement: Dr. Gregg Hageman
CDNA Library preparation: Dr. M. Bento Soares, Univeristy of lowa CDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa CDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa CDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa CDNA Eighting Library Esearchers may obtain clones from Research Genetics (www.resgen.com).
The following repetitive elements were found in this CDNA sequence: 1-22, AT_rich#Low_complexity (matched compliment)
Seg primer: M13 Forward
POLYA=Yes.
                                                                                                                                                                                                                                                                                                                                                                  BM664187 linear EST 27-FEB-2002
UI-E-CI1-age-d-10-0-UI.s1 UI-E-CI1 Homo sapiens cDNA clone
UI-E-CI1-age-d-10-0-UI 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bass 1 to 447)
Bonaldo, M.F., Lennon, G. and Soares, M.B.
Normalization and subtraction: two approaches to facilitate gene
             100.0%; Score 20; DB 9; Length 432; 100.0%; Pred. No. 27; ive 0; Mismatches 0; Indels
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/db xref="taxon:9606"
/clone="UI-E-CI1-age-d-10-0-UI"
/clone_lib="UI-E-CI1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TAG_LIB=UI-E-CI1
TAG_TISSUE=RPE and Choroid
TAG_SEQ=ACCTA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genome Rês. 6 (9), 791-806 (1996)
97044477
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                                                                                                                                             1 TTTCAACCAGATGGTCATTC
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                                                                                 20; Conservative
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             Query Match
Best Local Similarity
Matches 20; Conserv
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AUTHORS
TITLE
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I Unpublished (1997)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov

Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael

Email: cgapbs-remail.nih.gov

Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael

Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llh.gov/bbrp/image/image.html

Insert Length: 1419 Std Error: 0.00

Seq primer: -40mi3 fwd. ET from Amersham.

Location/Qualifiers
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/db_xref="taxon:9606"
/clone=!nbes:158466"
/clone=lib="NCI_CGAP_GG3"
/tissue_type="pooled germ cell tumors"
/lab.host="DH108"
/note="Vector: pT7T3D-Pac (Pharmacia) with a modified
polylinker; lst strand cDNA was prepared from 3 pooled
germ cell tumors, and was then primed with a Not I -
oligo(dT) primer. Double-stranded cDNA was ligated to Eco
RI adaptors (Pharmacia), digested with Not I and cloned
into the Not I and Eco RI sites of the modified pT7T3
vector. Library is not normalized. Library was
constructed by Bento Soares and M. Fatima Bonaldo. "
3716-3720, 3733-3735 (IMAGE CloneIDs 1257096-1258631, 1469064-1470883, 1475522-1476743); NCI CGAP Pr22 pool 1 LLAM 2457-2459, 2758-2759, 3062-306 (TWAGE CloneIDs 98608-986759, 1101192-1101959, 1217928-1220615); NCI CGAP_CO10 pool 1 LLAM 2644-2653, 2871-2872 (IMAGE CloneIDs 1057416-1061255, 1144584-1145351). Subtraction was performed as previously described (Bonaldo, Lennon & Soares (1996): Normalization and Subtraction: Two Approaches To Facilitate Gene Discovery. Genome Research 6, 791-806.
TAG LIBENCI CGAP GC4
TAG_TISSUE=Gene Cell
TAG_COAPARTC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   432 bp mRNA linear EST 27-AUG-1998 or41gl0.s1 NCI CGAP GC3 Homo sapiens CDNA clone IMAGE:1598466 3' similar to gb: \overline{L}0518\overline{6} FOCAL ADHESION KINASE (HUMAN); mRNA sequence. AA983514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EST 27-AUG-1998
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 20; DB 10; Length 429; 100.0%; Pred. No. 27; cive 0; Mismatches 0; Indels 0
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Matches 20; Conservative
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SOURCE
ORGANISM
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N42230/c
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AUTHORS
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                193
                                                                                             Local
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              TTTCAACCAGATGGTCATTC
                                            TTTCAACCAGATGGTCATTC 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 High quality sequence stops: 330 Source: IMAGE Consortium, LLNL free through LLNL; contact the This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1 Tel: 314 286 1800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J. Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,B., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.
The WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N42230 462 bp mRNA linear EST 24-JAN-19:
yw71h06.rl Soares_placenta_8to9weeks_2NbHP8to9W Homo sapiens cDNA
clone_IMAGE.257723 5' similar to gb:L05186 FOCAL ADHESION KINASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1995)
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hillier, L., Clark, N.,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                             Similarity
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Location/Qualifiers
                                                                              Conservative
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                                                                                                                                                        double-stranded cDNA was Size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library constructed by Bento Soares and M.Fatima Bonaldo."

1 96 c 89 g 139 t 1 others
                                                                                                                                                                                                                                                       /clone_lib="Soares_placenta_8to9weeks_2NbHP8to9W"
/dev_stage="two placentae: one from 8 weeks and another
from 9 weeks post conception"
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                                                                                                                                                                                                                                                                                                                                                                                      /db_xref="GDB:3887333"
/db_xref="taxon:9606"
/clone="IMAGE:257723"
                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Homo sapiens"
                                                                            100.0%; E
100.0%; E
tive 0;
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                                                                            Score 20; DB 1
Pred. No. 28;
0; Mismatches
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Pred. No. 27;
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                                              VERSION
KEYWORDS
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AUTHORS
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                                                                                                                                                                                                                           TTTCAACCAGATGGTCATTC 111
                                                                                                                                                                                                                                                         TTTCAACCAGATGGTCATTC 20
Homo sapiens
Eukaryota; Metazoa; (
Mammalia; Eutheria; I
1 (bases 1 to 492)
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similar to
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Unpublished (1997)
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Mammalia; Eutheria;
                                                                human.
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PM2-DT0042-100900-003-g03
BF084847
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Insert Length: 686 Std Error: 0.00
Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D.,
R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: M. Bento Soares, Ph.D.,
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AI341506.1 GI:4078433
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site 1: Not I; Site 2: Ecc RI; Plasmid DNA from the normalized library NCI_CGAP_GC4 was prepared, and se circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1257096-128631, 1469064-1470983, and 1475592-1476743). Subtraction by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="IMAGE:2009950"
/clone lib="NCI CGAP GC6"
/tissue_type="pooled germ cell tumors"
lab_host="DH10B"
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NCI CGAP GC6 Homo sapiens cDNA clone IMAGE:2009950 3'
gb:L05186 FOCAL ADHESION KINASE (HUMAN);, mRNA sequence.
                                                                                                                             GI:10878677
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Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                            Fax: +55-11-2707001

Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
This sequence was derived from the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=&t2=PM2-DT0042-100
900-003-903&E3=2000-09-10&E4=1)
Seq primer: puc 18 forward
High quality sequence start: 5
High quality sequence stop: 492.
Location/Qualifiers
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/db_xref="taxon:9606"

/db_xref="taxon:9606"

/dcloe_lib="byr0042"

/dcv stage="Adult"

/note="Organ: denis drash; Vector: pucl8; Site_1: Smal;

Site_2: Smal; A mini-library was made by cloning products

derived from ORESTES PCR (U.S. Letters Patent application

No. 196,716 - Ludwig Institute for Cancer Research)

profiles into the pUC 18 vector. Reverse transcription of
tissue mRNA and cDNA amplification were performed under
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 492)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Brunscein,A., deoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., de Oliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and Simpson, A.J.
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                                                                                                                                                                                                                                               Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
Brazil
                                                                                                                                Shotgun sequencing of the human transcriptome with ORF expressed
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Proc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)
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Laboratory of Cancer Genetics
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/organism="Homo sapiens"

/db xref="taxon:966"

/clon=lib="DT0042"

/dev stage="Adult"

/note="Organ: denis drash; Vector: puc18; Site_1: Smal;

Site_2: Smal; A mini-library was made by cloning products
derived from ORESTES PCR (U.S. Letters Patent application
No. 196,716 - Ludwig Institute for Cancer Research)
profiles into the pUC 18 vector. Reverse transcription of
tissue mRNA and cDNA amplification were performed under
low stringency conditions."
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Email: cgapbs-r@mail.nih.gov
cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima Bonaldo
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DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-GGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 752 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stope: 344.
                                                                                                                                                              Fax: +55-11-2707001
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=PM2&t2=PM2-DT0042-
190201-005-903&t3=2001-02-19&t4=1)
Seq primer: puc 18 forward
High quality sequence start: 5
High quality sequence stop: 492.
Location/Qualifiers
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1 (bases 1 to 527)
Nati-Caph http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index
Unpublished (1997)
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
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/db_xref="taxon:9606"
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BASE COUNT
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Search completed: January 12, 2003, 11:09:54 Job time : 1997 secs
                                                          Query Match 100.0%; Score 20; DB 9; Length 527; Best Local Similarity 100.0%; Pred. No. 30; Matches 20; Conservative 0; Mismatches 0; Indels
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